

7 Anhang

7.1 Software

Betriebssysteme:

- NT
- OS/2

Anwendungssoftware:

- OPUS-Ident (Bruker)
Spektroskopie-Software mit „IDENT“-Erweiterung zur Erstellung von Spektrenbibliotheken
- Visual Studio 97 (Microsoft)
Programmiersoftware, welche u.a. C++ umfaßt
- Matlab, Wavelet-toolbox (The Math Works Inc.)
Software mit umfangreichen mathematischen Funktionen und Wavelet-Erweiterung zur Wavelet-Transformation mit Daubechies-Wavelets
- Specwork (C) 1997 von Dr. U. Depczynski,
Software für die Wavelet-Transformation mit S-Wavelets

7.2 Substanzliste

| | | | | | |
|---------|---------|--|----|--------|--|
| Lfd.Nr. | Art.Nr. | Artikelname | 45 | 101187 | Ammoniumnitrat reinst |
| 1 | 100010 | Acetanilid reinst krist | 46 | 101190 | Di-Ammoniumoxalat-Monohydrat |
| 2 | 100085 | Acetylsalicylsäure | 47 | 101200 | Ammoniumperoxodisulfat reinst |
| 3 | 100090 | Adipinsäure reinst | 48 | 101208 | Ammoniumthiosulfat LAB |
| 4 | 100102 | 4-Aminobenzoesäure reinst | 49 | 101211 | Ammoniumsulfat f. b. Zw. |
| 5 | 100103 | Amidoschwefelsäure z.A. | 50 | 101212 | Ammoniumthiocyanat reinst |
| 6 | 100129 | L-Asparaginsäure reinst | 51 | 101218 | di-Ammoniumhydrogenphosphat |
| 7 | 100130 | Benzoessäure gepulvert | | | m. Fließmittel |
| 8 | 100138 | 3,5-Dinitrobenzoessäure | 52 | 101226 | Ammoniummonovanadat z.A. |
| 9 | 100162 | Borsäure gepulvert reinst | 53 | 101407 | Kristallviolett C.I. 42555 |
| 10 | 100163 | Di-Bortrioxid | 54 | 101492 | L(+)-Araboise f. Mikrobiologie |
| 11 | 100241 | Citronensäure wasserfrei | 55 | 101543 | L-Argininmonohydrochlorid |
| 12 | 100242 | Citronensäure Monohydrat krist | 56 | 101565 | L-Asparagin-Monohydrat reinst |
| 13 | 100243 | Citronensäure Monohydrat gepulvert | 57 | 101719 | Bariumchlorid-Dihydrat z.A. |
| 14 | 100289 | L-Glutamin f. bioch. Zwecke | 58 | 101798 | 1H-Benzotriazol techn. |
| 15 | 100290 | L-Glutaminsäure reinst | 59 | 101838 | Curco BS |
| 16 | 100383 | DL-Äpfelsäure, Lebensmittelqualität | 60 | 101841 | Oxalsäure-bis-(Cyclohexylidenhydrazid) z.A. |
| 17 | 100492 | Oxalsäure-Dihydrat reinst | 61 | 102070 | Calciumchlorid 90-95% entwässert |
| 18 | 100623 | Pikrinsäure z.A. (0,5ml Wasser/g) | 62 | 102071 | Calciumcarbonat f. op. Glas |
| 19 | 100631 | Salicylsäure reinst | 63 | 102072 | Calciumchlorid-Hexahydrat reinst |
| 20 | 100662 | Sorbinsäure | 64 | 102102 | Calciumlactat, löslich reinst |
| 21 | 100671 | Stearinsäure | 65 | 102110 | Calciumhydroxid FCC Lebensmittelqualität |
| 22 | 100668 | Bernsteinsäure z.A. | 66 | 102127 | Beta-Cyclodextrin |
| 23 | 100773 | Tannin gepulvert rein | 67 | 102144 | Calciumhydrogenphosphat wasserfrei |
| 24 | 100802 | L+Weinsäure krist | 68 | 102146 | Calciumhydrogenphosphat Dihydrat |
| 25 | 100807 | Trichloressigsäure | 69 | 102148 | Calcium-D-Saccharat reinst |
| 26 | 100846 | Adonit (Ribit) f.d. Mikrobiol. | 70 | 102154 | Calciumstearat |
| 27 | 100848 | Aesculin rein | 71 | 102162 | Calciumsulfat Hemihydrat gebrannt |
| 28 | 100877 | Ethyl-4-aminobenzoat (Benzocain), reinst | 72 | 102166 | D(-)-Campher raffiniert, gepulvert |
| 29 | 100886 | Ethyl-4-hydroxybenzoat Natriumsalz | 73 | 102196 | Hydroxylapatit f. Biokeramik |
| 30 | 100887 | Ethyl-4-hydroxybenzoat | 74 | 102352 | Cellobiose f. bioch. Zwecke |
| 31 | 100944 | Etylendinitrilotetra-Essigsäure reinst | 75 | 102424 | Chloramin T Trihydrat reinst |
| 32 | 100963 | DL-Alanin für bioch. Zwecke | 76 | 102425 | Chloralhydrat |
| 33 | 100989 | Cetylalkohol (1-Hexadecanol) reinst | 77 | 102507 | Cinchoninhydrochlorid Monohydrat |
| 34 | 101005 | 1,1,1-Trichlor-2-Methyl-2-Propanol-Hemihydrat (Chlorbutanol) | 78 | 102530 | Cobalt-II-acetat Tetrahydrat |
| 35 | 101072 | Aluminiumchloridhydroxid-Allantoin | 79 | 102584 | Coffein rein |
| 36 | 101091 | Aluminiumhydroxid gepulvert | 80 | 102836 | L-Cystin reinst |
| 37 | 101107 | Aluminiumstearat | 81 | 102838 | L-Cystein für bioch. Zwecke |
| 38 | 101124 | Ammoniumdihydrogenphosphat reinst krist. | 82 | 102839 | L-Cysteinhydrochlorid-Monohydrat für bioch. Zwecke |
| 39 | 101125 | Ammoniumbromid reinst | 83 | 102851 | Calciumlaevulinat reinst |
| 40 | 101131 | Ammoniumhydrogencarbonat | 84 | 102968 | Neocuproin (2,9-Dimethyl-1,10-Phenanthrolin) z.A. |
| 41 | 101141 | Ammoniumchlorid rein | 85 | 103006 | Dextrin weiß |
| 42 | 101149 | Ammoniumdichromat reinst krist | 86 | 103149 | (+)-Ephedrinhydrochlorid |
| 43 | 101155 | Di-Ammoniumhydrogencitrat reinst | 87 | 103672 | Colesterin gepulvert |
| 44 | 101182 | Ammoniumheptamolybdat-Tetrahydrat z.A. | 88 | 103767 | Ammoniumformiat reinst |
| | | | 89 | 103883 | Eisen-III-nitrat Nonahydrat z.A. |
| | | | 90 | 103929 | Fluorescein-Natrium reinst |
| | | | 91 | 103962 | Eisen(II)-fumarat, reinst |
| | | | 92 | 104005 | Paraformaldehyd reinst |

| | | | | | |
|-----|--------|--|-----|--------|--|
| 93 | 104061 | D(+)-Galactose reinst | 139 | 106129 | 3-Morpholinopropansulfonsäure |
| 94 | 104078 | Gelatine gepulvert | 140 | 106223 | 1-Naphthol |
| 95 | 104112 | Calciumglycerophosphathydrat | 141 | 106247 | Naphazolinnitrat, reinst |
| 96 | 104128 | Auxillin (Gibberillin mit etwa 90% wirks. Gibberillin A3) | 142 | 106249 | Naphazolinhydrochlorid, reinst |
| 97 | 104168 | Beta-Glycerophosphat Dinatrium- salz-Pentahydrat | 143 | 106346 | Natriumdihydrogenphosphat- Monohydrat z.A. |
| 98 | 104219 | Guanidiniumchlorid f. b. Zw. | 144 | 106371 | Natriumborhydrid z.A. |
| 99 | 104228 | Gummi Arabisch, sprühgetrocknet | 145 | 106392 | Natriumcarbonat wasserfrei |
| 100 | 104343 | Methenamin | 146 | 106431 | Tri-Natriumcitrat-5,5-Hydrat reinst |
| 101 | 104349 | Homatropinmethylbromid | 147 | 106432 | Tri-Natriumcitrat-Dihydrat |
| 102 | 104350 | L-Histidinmonohydrochlorid- Monohydrat f. b. Zw. | 148 | 106445 | Natrium-L-glutamat-Monohydrat |
| 103 | 104351 | L-Histidin f. b. Zw. | 149 | 106446 | tri-Natriumcitrat Dihydrat gepul- vert reinst |
| 104 | 104370 | Histamindihydrochlorid reinst | 150 | 106467 | Natriumhydroxid granuliert |
| 105 | 104506 | L-Hydroxyprolin f. b. Zw. | 151 | 106544 | Natriumnitrit reinst |
| 106 | 104541 | Dragiersuspension f. farbint. Dra- gees | 152 | 106600 | Natriumsalicylat krist |
| 107 | 104610 | Hydrochinon Fotopur | 153 | 106619 | Brenztraubensäure-Na.-Salz f. bioch. Zwecke |
| 108 | 104731 | Myo-Inosit | 154 | 106663 | Di-Natriumtartrat-Dihydrat z. A. |
| 109 | 104751 | Jodoform gepulvert reinst | 155 | 106688 | Natriumazid reinst |
| 110 | 104819 | Kalium-Titriplex Dihydrat z.A. | 156 | 106756 | Methyl-4-hydroxybenzoat Natri- umsalz |
| 111 | 104820 | Kaliumacetat reinst | 157 | 106757 | Methyl-4-hydroxybenzoat |
| 112 | 104892 | Kaliumhydrogentartrat reinst | 158 | 106832 | Natriumcyclamat |
| 113 | 104956 | Tri-Kaliumcitrat-Monohydrat reinst | 159 | 107069 | Phloroglucin (1,3,5- Trihydroxybenzol) |
| 114 | 104982 | Kaliumhexacyanoferrat-II- Trihydrat reinst | 160 | 107225 | 1,10-Phenanthrolin-Monohydrat z.A. |
| 115 | 105061 | Kaliumnitrat reinst | 161 | 107230 | Phenolphthalein |
| 116 | 105072 | Di-Kaliumoxalat-Monohydrat reinst | 162 | 107256 | L-Penylalanin f. b. Zw. |
| 117 | 105099 | Di-Kaliumhydrogenphosphat- Trihydrat z.AAv10 | 163 | 107260 | Phenylsalicylat |
| 118 | 105101 | Di-Kaliumhydrogenphosphat was- serfrei reinst | 164 | 107311 | Pilocarpinhydrochlorid, krist reinst |
| 119 | 105102 | Tri-Kaliumphosphat-Trihydrat reinst | 165 | 107427 | Propyl-4-Hydroxybenzoat |
| 120 | 105118 | Kaliumsorbit Pulver | 166 | 107428 | Propyl-4-hydroxybenzoat Natrium- salz |
| 121 | 105125 | Kaliumthiocyanat z.A. | 167 | 107434 | L-Prolin f. b. Zwecke |
| 122 | 105321 | D(-)-Fructose reinst | 168 | 107467 | Procainhydrochlorid reinst |
| 123 | 105360 | L-Leucin | 169 | 107549 | Raffinose-Pentahydrat (Melitose) |
| 124 | 105362 | L-Isoleucin | 170 | 107590 | Resorcin reinst |
| 125 | 105645 | DL-alpha-Liponsäure, reinst | 171 | 107605 | D(-)-Ribose |
| 126 | 105683 | Trilithiumcitrat-Tetrahydrat | 172 | 107606 | 2-desoxy-D-Ribose |
| 127 | 105691 | Lithiumhydroxid | 173 | 107640 | Rübenzucker, Raffinade |
| 128 | 105700 | L-Lysinmonohydrochlorid f. b. Zw. | 174 | 107656 | Lactose-Monohydrat gepulvert |
| 129 | 105707 | L-Methionin | 175 | 107665 | Salicin (2-O-(B-D- Glucopyranosido)-benzylalkohol) |
| 130 | 105870 | Magnesiumhydroxid, reinst | 176 | 107682 | Seife medizinisch |
| 131 | 105873 | Magnesiumperchlorat-Hydrat (ca. 83%-ig) | 177 | 107769 | L-Serin |
| 132 | 105876 | Magnesiumstearat gefällt fein ge- pulvert | 178 | 107959 | Süßholzextrakt sprühgetrocknet |
| 133 | 105904 | Magnesiumcitrat, reinst | 179 | 107978 | Thioharnstoff |
| 134 | 105911 | Maltose Monohydrat, krist | 180 | 108035 | Sulfanilamid reinst |
| 135 | 105924 | Mangan-II-carbonat-Hydrat z. A. | 181 | 108042 | Saccharin Natrium |
| 136 | 105984 | D(+)-Mannose | 182 | 108085 | Kaliumnatriumtartrat-Tetrahydrat krist reinst |
| 137 | 105995 | (-)-Menthol krist | 183 | 108092 | Kaliumantimon-(III)-oxid-Tartrat- Hemihydrat reinst |
| 138 | 106126 | 2-Morpholinoethansulfonsäure- Monohydrat | 184 | 108170 | Thioacetamid z.A. |
| | | | 185 | 108196 | Tablettierhilfsmittel D |
| | | | 186 | 108197 | Tablettierhilfsmittel K |
| | | | 187 | 108198 | Tablettierhilfsmittel T |
| | | | 188 | 108337 | D+Glucose wasserfrei |

| | | | | | |
|-----|--------|--|-----|--------|---|
| 189 | 108346 | D+Glucose Monohydrat | 223 | 115419 | Cetylpalmitat |
| 190 | 108353 | Trehalose | 224 | 115940 | Kristallviolett C.I. 42555 |
| 191 | 108371 | L-Tyrosin f. b. Zw. | 225 | 118304 | Pentan-1-Sulfonsäurenatriumsalz |
| 192 | 108374 | L-Tryptophan f. b. Zw. | 226 | 118305 | Hexan-1-sulfonsäure Natriumsalz |
| 193 | 108382 | Tris(hydroxymethyl)-aminomethan (Trometamol) | 227 | 118306 | Heptan-1-sulfonsäure Natriumsalz |
| 194 | 108411 | L-Threonin f. b. Zw. | 228 | 118307 | Octan-1-sulfonsäure Natriumsalz |
| 195 | 108418 | Titriplex III, (Ethylendinitrilote- traessigsäure Dinatriumsalz Dihydrat) | 229 | 118308 | Decan-1-sulfonsäure Natriumsalz |
| 196 | 108439 | Calcium-Titriplex Dihydrat | 230 | 124511 | 1,4-Dithioerythrit für bioch. Zwek- ke |
| 197 | 108484 | Harnstoff, Perlform reinst | 231 | 158626 | Tri-Lithiumcitrat ca. 2 mol Wasser |
| 198 | 108486 | Harnstoff reinst krist | 232 | 159605 | 4-Aminophenol |
| 199 | 108495 | L-Valin | 233 | 270994 | Thiaminchloridhydrochlorid |
| 200 | 108510 | Vanillin, Lebensmittelqual. | 234 | 271436 | Adenosin-5-diphosphorsäure Monokaliumsalz Dihydrat |
| 201 | 108865 | Zinkstearat | 235 | 277353 | Photo Rex |
| 202 | 110110 | 2-[4-(2-Hydroxyethyl)-1- piperazinyl]-ethansulfonsäure | 236 | 278343 | Dextroseandyrid |
| 203 | 110220 | Piperazin-1,4- bis(ethansulfonsäure) | 237 | 450890 | Saponin |
| 204 | 110550 | Creatinphosphorsäure Dinatrium- salz-Tetrahydrat | 238 | 451318 | DL-Methionin |
| 205 | 110783 | Di-Lithiumtetraborat (Spectromelt A 10) | 239 | 458038 | Ethylcellulose |
| 206 | 110887 | Dichlorisocyanursäure Natriumsalz Dihydrat | 240 | 458162 | Aerosol OT |
| 207 | 111037 | Trinatriumcitrat | 241 | 458268 | Saponin |
| 208 | 111352 | Molinon Instant | 242 | 500010 | Folsäure rein |
| 209 | 111452 | Phthaldialdehyd | 243 | 500013 | Rutin (Rutosid) |
| 210 | 111474 | 1,4-Dithiothreit | 244 | 500090 | L(+)-Ascorbylpalmitat |
| 211 | 111685 | Stärke (aus Weizen) | 245 | 500115 | Cholinhydrogentartrat, Lebens- mittelqualität |
| 212 | 112081 | S-Butyrylthiocholiodid f. Bioch. Zwecke | 246 | 500117 | Cholinchlorid reinst |
| 213 | 112240 | Melibiose Monohydrat | 247 | 500157 | Calcium-D(+)-Pantothenat |
| 214 | 112422 | N-Acetyl-L-cystein | 248 | 500190 | Glycin krist |
| 215 | 112428 | Amygdalin f. Bioch. Zwecke | 249 | 500224 | Pyridoxolhydrochlorid |
| 216 | 112488 | N-Acetyl-DL-tryptophan reinst en- dotoxiarm | 250 | 500257 | Riboflavin |
| 217 | 113125 | 2,6-Dibromchinon-4-chlorimid (max. 30% Wasser) z.A. | 251 | 500259 | Riboflavin-5-phosphat Mononatri- umsalz Dihydrat |
| 218 | 114266 | Polyvinylalkohol | 252 | 500297 | Nicotinsäureamid |
| 219 | 115226 | N-(2-Acetamido)-2- aminoethansulfonsäure | 253 | 500675 | Thiamindisulfid |
| 220 | 115230 | 3-[4-(2-Hydroxyethyl)-1- piperazinyl]-ropansulfonsäure | 254 | 500724 | Vitamin A-acetat Trockenpulver |
| 221 | 115231 | 2-[4-(2-Hydroxyethyl)-1- piperazinyl]-ethansulfonsäure Natriumsalz | 255 | 500859 | Vitamin E-Trockenpulver (50g DL-A-Tocopherolacetat in 100g Substanz) |
| 222 | 115299 | Dextromethorphanhydrobromid | 256 | 500980 | Thiaminnitrat (Vitamin B1 Mono- nitrat) |
| | | | 257 | 501260 | Pyridoxol |
| | | | 258 | 501551 | Vitamin D3-Trockenpulver |
| | | | 259 | 501606 | D+Biotin-konz. 2% gespr. |
| | | | 260 | 501902 | Trihydroxyethylrutin |
| | | | 261 | 524950 | Vitamin B12 |
| | | | 262 | 820927 | Octansäure Natriumsalz |
| | | | 263 | 907327 | Piperazin-Hexahydrat |

7.3 Validierungsreports

NIR (Standardmethode, Vektornormierung und 1.Ableitung)

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Method file:                E:\IDL\IB\IPI\KL1\MTHD3\SND19d.FAA
from (date):                4/ 1/99
  (time):                   19:11:4
Algorithm:                  Standard
For Threshold Individual:   Nein
No. of used factor sp.:    -
For Threshold Info Entry:  1
No. of Hits to be Listed:  10
Vector normalized spectra: Ja
No. of Spectra:            263
Delta x of x raster:       3.857
FXV (or LXV) mod Delta x:  7.4607e-014
min x of x raster:         3999.71
max. x of x raster:        11999.1
Whole x range:             Ja
Nr. of x Points in all x Ranges: 2075
X-Ranges:                  1
From:                      3999.71
  to:                      11999.1
  Weight:                  1
  Reprolevel:              1
BlockID:                   4111
Order of Derivative:       0
Description:
For Threshold Info Entry:  TEXT:Compound Name
Path of origin reference spectra: E:\EINZEL\KL1
Smoothing points:          1
Method Released:           0
Order of Internal Derivation: 1
Smoothing Points for Internal Derivation: 9
Reduction Factor:          1
Use Thresh. From Info:     Nein
Use Thresh. From Info Entry: 1
Use Thresh. From Info Entry: TEXT:Compound Name

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Check Average Spectra

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Constant conf. level : Fixed algorithm, Single Sphere Method

Confidence Region = 2 x Threshold

Spectra, which can be confused with other references:

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 100242.1 | 100242 Citronensäure Monohydr | 1.004088 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 100243.1 | 100243 Citronensäure Monohydr | 0.925198 | 0.503104 |
| 100802.1 | 100802 L+Weinsäure krist Av | 0.947264 | 0.587235 |
| 104219.1 | 104219 Guanidiniumchlorid f. | 0.963606 | 0.220449 |
| 105099.1 | 105099 Di-Kaliumhydrogenphosp | 0.906586 | 0.484008 |
| 907327.1 | 907327 Piperazin-Hexahydrat | 0.937345 | 1.040533 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 101124.1 | 101124 Ammoniumdihydrogenphos | 0.758398 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 101131.1 | 101131 Ammoniumhydrogencarbon | 0.680011 | 0.773526 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 101131.1 | 101131 Ammoniumhydrogencarbon | 0.773526 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 101124.1 | 101124 Ammoniumdihydrogenphos | 0.680011 | 0.758398 |
| 101218.1 | 101218 di-Ammoniumhydrogenpho | 0.670188 | 0.956888 |

| File Name | Sample Name | Confidence Region |
|-------------------|----------------------------------|-------------------|
| 101187.1 | 101187 Ammoniumnitrat reinst | 0.391632 |
| ----- | | |
| Overlapping with: | | Hit |
| 101211.1 | 101211 Ammoniumsulfat f. b. Z | 0.251700 0.448471 |
| File Name | Sample Name | Confidence Region |
| 101200.1 | 101200 Ammoniumperoxodisulfat | 0.471006 |
| ----- | | |
| Overlapping with: | | Hit |
| 101187.1 | 101187 Ammoniumnitrat reinst | 0.420118 0.391632 |
| 101211.1 | 101211 Ammoniumsulfat f. b. Z | 0.461298 0.448471 |
| File Name | Sample Name | Confidence Region |
| 101208.1 | 101208 Ammoniumthiosulfat LA | 0.532495 |
| ----- | | |
| Overlapping with: | | Hit |
| 101211.1 | 101211 Ammoniumsulfat f. b. Z | 0.521505 0.448471 |
| File Name | Sample Name | Confidence Region |
| 101211.1 | 101211 Ammoniumsulfat f. b. Z | 0.448471 |
| ----- | | |
| Overlapping with: | | Hit |
| 101187.1 | 101187 Ammoniumnitrat reinst | 0.251700 0.391632 |
| File Name | Sample Name | Confidence Region |
| 101218.1 | 101218 di-Ammoniumhydrogenpho | 0.956888 |
| ----- | | |
| Overlapping with: | | Hit |
| 100807.1 | 100807 Trichloressigsäure A | 0.858144 0.624967 |
| 101124.1 | 101124 Ammoniumdihydrogenphos | 0.884729 0.758398 |
| 101131.1 | 101131 Ammoniumhydrogencarbon | 0.670188 0.773526 |
| 103767.1 | 103767 Ammoniumformiat reinst | 0.884624 0.539182 |
| File Name | Sample Name | Confidence Region |
| 101407.1 | 101407 Kristallviolett C.I. 42 | 0.704777 |
| ----- | | |
| Overlapping with: | | Hit |
| 115940.1 | 115940 Kristallviolett C.I. 42 | 0.028305 0.333492 |
| File Name | Sample Name | Confidence Region |
| 102072.1 | 102072 Calciumchlorid-Hexahyd | 1.134704 |
| ----- | | |
| Overlapping with: | | Hit |
| 100492.1 | 100492 Oxalsäure-Dihydrat rei | 0.878870 0.396618 |
| 100623.1 | 100623 Pikrinsäure z.A. (0,5m | 1.131745 0.350166 |
| 101719.1 | 101719 Bariumchlorid-Dihydrat | 1.126216 0.314242 |
| 102127.1 | 102127 Beta-Cyclodextrin Av | 1.109518 0.288776 |
| 102144.1 | 102144 Calciumhydrogenphospha | 1.074647 0.531808 |
| 103006.1 | 103006 Dextrin weiß Av. of | 1.118213 0.090826 |
| 103883.1 | 103883 Eisen-III-nitratNonahy | 1.074546 0.210122 |
| 104228.1 | 104228 Gummi Arabisch, sprühg | 1.110395 0.099201 |
| 104982.1 | 104982 Kaliumhexacyanoferrat- | 1.123569 0.476212 |
| 105099.1 | 105099 Di-Kaliumhydrogenphosp | 1.104069 0.484008 |
| 105924.1 | 105924 Mangan-II-carbonat-Hyd | 1.102652 0.238062 |
| 106346.1 | 106346 Natriumdihydrogenphosp | 1.040737 0.618181 |
| 108484.1 | 108484 Harnstoff, Perlform re | 1.110541 0.174455 |
| 501606.1 | 501606 D+Biotin-konz. 2% gespr. | 1.121857 0.147494 |
| 907327.1 | 907327 Piperazin-Hexahydrat | 1.053368 1.040533 |
| File Name | Sample Name | Confidence Region |
| 104350.1 | 104350 L-Histidinmonhydrochl | 1.037221 |
| ----- | | |
| Overlapping with: | | Hit |
| 524950.1 | 524950 Vitamin B12 Av. of 1 | 0.984871 0.462871 |
| File Name | Sample Name | Confidence Region |
| 105700.1 | 105700 L-Lysinmonhydrochlori | 0.912729 |
| ----- | | |
| Overlapping with: | | Hit |
| 108495.1 | 108495 L-Valin Av. of 12 | 0.832032 0.220122 |
| File Name | Sample Name | Confidence Region |

| | | | | | |
|-------------------|-------------|------------------------|----------|------------|----------|
| 105995.1 | 105995 | (-)-Menthol krist | Av | | 0.922771 |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 102166.1 | 102166 | D(-)-Campher raffinier | | 0.894854 | 0.590237 |
| 103672.1 | 103672 | Colesterin gepulvert | | 0.786379 | 0.068346 |
| 115419.1 | 115419 | Cetylpalmitat | Av. of | 0.898198 | 0.847277 |
| 118304.1 | 118304 | Pentan-1-Sulfonsäurena | | 0.917566 | 0.579100 |
| 458162.1 | 458162 | Aerosol OT | Av. of 7 | 0.801861 | 0.650112 |
| 500859.1 | 500859 | Vitamin E-Trockenpulve | | 0.905776 | 0.069755 |
| File Name | Sample Name | | | Confidence | Region |
| 106432.1 | 106432 | Tri-Natriumcitrat-Dihy | | 0.220126 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 106446.1 | 106446 | tri-Natriumcitrat Dihy | | 0.068439 | 0.156468 |
| File Name | Sample Name | | | Confidence | Region |
| 106446.1 | 106446 | tri-Natriumcitrat Dihy | | 0.156468 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 106432.1 | 106432 | Tri-Natriumcitrat-Dihy | | 0.068439 | 0.220126 |
| File Name | Sample Name | | | Confidence | Region |
| 107428.1 | 107428 | Propyl-4-hydroxybenzoa | | 0.979749 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 100886.1 | 100886 | Ethyl-4-hydroxybenzoat | | 0.813607 | 0.309095 |
| 107427.1 | 107427 | Propyl-4-Hydroxybenzoa | | 0.877347 | 0.205656 |
| 458038.1 | 458038 | Ethylcellulose | Av. o | 0.924451 | 0.138814 |
| File Name | Sample Name | | | Confidence | Region |
| 107682.1 | 107682 | Seife medizinisch | Av | 0.265627 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 105876.1 | 105876 | Magnesiumstearat gefäl | | 0.247753 | 0.080412 |
| File Name | Sample Name | | | Confidence | Region |
| 111352.1 | 111352 | Molinon Instant | Av. | 0.635889 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 458268.1 | 458268 | Saponin | Av. of 5 | 0.607193 | 0.393725 |
| File Name | Sample Name | | | Confidence | Region |
| 111685.1 | 111685 | Stärke (aus Weizen) | | 0.106977 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 103006.1 | 103006 | Dextrin weiß | Av. of | 0.099303 | 0.090826 |
| File Name | Sample Name | | | Confidence | Region |
| 115419.1 | 115419 | Cetylpalmitat | Av. of | 0.847277 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 100989.1 | 100989 | Cetylalkohol (1-Hexade | | 0.702835 | 0.391486 |
| 458162.1 | 458162 | Aerosol OT | Av. of 7 | 0.819149 | 0.650112 |
| File Name | Sample Name | | | Confidence | Region |
| 115940.1 | 115940 | Kristallviolet C.I. 42 | | 0.333492 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 101407.1 | 101407 | Kristallviolet C.I. 42 | | 0.028305 | 0.704777 |
| File Name | Sample Name | | | Confidence | Region |
| 118304.1 | 118304 | Pentan-1-Sulfonsäurena | | 0.579100 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 118305.1 | 118305 | Hexan-1-sulfonsäure Na | | 0.396403 | 0.583429 |
| 118306.1 | 118306 | Heptan-1-sulfonsäure N | | 0.467386 | 0.459277 |
| File Name | Sample Name | | | Confidence | Region |
| 118305.1 | 118305 | Hexan-1-sulfonsäure Na | | 0.583429 | |
| ----- | | | | | |
| Overlapping with: | | | | Hit | |
| 118304.1 | 118304 | Pentan-1-Sulfonsäurena | | 0.396403 | 0.579100 |
| 118306.1 | 118306 | Heptan-1-sulfonsäure N | | 0.242794 | 0.459277 |

| | | | | |
|-------------------|-------------|------------------------|------------|----------|
| 118307.1 | 118307 | Octan-1-sulfonsäure Na | 0.464148 | 0.521464 |
| File Name | Sample Name | | Confidence | Region |
| 118306.1 | 118306 | Heptan-1-sulfonsäure N | 0.459277 | |
| Overlapping with: | | | Hit | |
| 118305.1 | 118305 | Hexan-1-sulfonsäure Na | 0.242794 | 0.583429 |
| 118307.1 | 118307 | Octan-1-sulfonsäure Na | 0.297615 | 0.521464 |
| File Name | Sample Name | | Confidence | Region |
| 118307.1 | 118307 | Octan-1-sulfonsäure Na | 0.521464 | |
| Overlapping with: | | | Hit | |
| 118305.1 | 118305 | Hexan-1-sulfonsäure Na | 0.464148 | 0.583429 |
| 118306.1 | 118306 | Heptan-1-sulfonsäure N | 0.297615 | 0.459277 |
| 118308.1 | 118308 | Decan-1-sulfonsäure Na | 0.317593 | 0.333075 |
| File Name | Sample Name | | Confidence | Region |
| 118308.1 | 118308 | Decan-1-sulfonsäure Na | 0.333075 | |
| Overlapping with: | | | Hit | |
| 118307.1 | 118307 | Octan-1-sulfonsäure Na | 0.317593 | 0.521464 |
| File Name | Sample Name | | Confidence | Region |
| 450890.1 | 450890 | Saponin Av. of 9 | 0.289127 | |
| Overlapping with: | | | Hit | |
| 107656.1 | 107656 | Lactose-Monohydrat gep | 0.203503 | 0.045294 |
| 501551.1 | 501551 | Vitamin D3-Trockenpulv | 0.174995 | 0.453883 |
| File Name | Sample Name | | Confidence | Region |
| 501551.1 | 501551 | Vitamin D3-Trockenpulv | 0.453883 | |
| Overlapping with: | | | Hit | |
| 107656.1 | 107656 | Lactose-Monohydrat gep | 0.253547 | 0.045294 |
| 450890.1 | 450890 | Saponin Av. of 9 | 0.174995 | 0.289127 |
| File Name | Sample Name | | Confidence | Region |
| 907327.1 | 907327 | Piperazin-Hexahydrat | 1.040533 | |
| Overlapping with: | | | Hit | |
| 100242.1 | 100242 | Citronensäure Monohydr | 0.937345 | 1.004088 |
| 100492.1 | 100492 | Oxalsäure-Dihydrat rei | 0.961665 | 0.396618 |
| 103883.1 | 103883 | Eisen-III-nitratNonahy | 0.914132 | 0.210122 |
| 105099.1 | 105099 | Di-Kaliumhydrogenphosp | 0.819565 | 0.484008 |
| 108484.1 | 108484 | Harnstoff, Perlform re | 0.988493 | 0.174455 |
| 108486.1 | 108486 | Harnstoff reinst krist | 1.018105 | 0.298561 |

Summary:

Average Spectra, which can be confused with other references: 29 of 263

Average Spectra, which can be uniquely identified: 234 of 263

| | | |
|----------|--------|-------------------------|
| 100010.1 | 100010 | Acetanilid reinst kris |
| 100085.1 | 100085 | Acetylsalicylsäure A |
| 100090.1 | 451552 | Adipinsäure reinst A |
| 100102.1 | 100102 | 4-Aminobenzoesäure rei |
| 100103.1 | 100103 | Amidoschwefelsäure z.A |
| 100126.1 | 100126 | L-Asparaginsäure für b |
| 100130.1 | 100130 | Benzoessäure gepulvert |
| 100138.1 | 100138 | 3,5-Dinitrobenzoessäure |
| 100162.1 | 100162 | Borsäure gepulvert rei |
| 100163.1 | 100163 | Di-Bortrioxid Av. of |

Raman (Standardmethode, Vektornormierung und 1.Ableitung)

```

Method file:                D:\IDLIB\IP1\KL2\MTHD3\SND19c.FAA
from (date):                5/ 1/99
  (time):                  15: 2:48
Algorithm:                  Standard
For Threshold Individual:   Nein
No. of used factor sp.:    -
For Threshold Info Entry:  1
No. of Hits to be Listed:  10
Vector normalized spectra: Ja
No. of Spectra:            263
Delta x of x raster:       0.96425
FXV (or LXV) mod Delta x:  0.280662
min x of x raster:         99.5984
max x of x raster:         3499.54
Whole x range:             Ja
No. of x Points in all x Ranges: 3527
X-Ranges:                  1
From:                       99.5984
  to:                       3499.54
  Weight:                   1
  Replevel:                 1
BlockID:                    10255
Order of Derivative:        0
Description:
For Threshold Info Entry:
Path of origin reference spectra: D:\EINZEL\KL21
Smoothing points:          1
Method Released:           0
Order of Internal Derivation: 1
Smoothing Points for Internal Derivation: 9
Reduction Factor:          1
Use Thresh. From Info:     Nein
Use Thresh. From Info Entry: 1
Use Thresh. From Info Entry:

```

Check Average Spectra

=====

Constant conf. level : Fixed algorithm, Single Sphere Method

Confidence Region = 2 x Threshold

Spectra, which can be confused with other references:

| File Name | Sample Name | Confidence Region |
|-----------|----------------------------------|-------------------|
| 100242.1 | 100242 Citronens,,ure Monohydr | 0.458916 |

| Overlapping with: | Hit | |
|-------------------|-----------------------------------|-------------------|
| 100243.1 | 100243 Citronens,,ure Monchhydr | 0.093363 0.544981 |

| File Name | Sample Name | Confidence Region |
|-----------|----------------------------------|-------------------|
| 100243.1 | 100243 Citronens,,ure Monohydr | 0.544981 |

| Overlapping with: | Hit | |
|-------------------|----------------------------------|-------------------|
| 100242.1 | 100242 Citronens,,ure Monohydr | 0.093363 0.458916 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 100989.1 | 100989 Cetylalkohol (1-Hexade | 0.589218 |

| Overlapping with: | Hit | |
|-------------------|----------------------------------|-------------------|
| 100671.1 | 100671 Stearins,,ure Av. of | 0.298740 0.026060 |
| 101107.1 | 101107 Aluminiumstearat Av. | 0.405257 0.075229 |
| 102154.1 | 102154 Calciumstearat Av. o | 0.424873 0.107628 |
| 105876.1 | 105876 Magnesiumstearat gef,,l | 0.431600 0.123470 |
| 107682.1 | 107682 Seife medizinisch Av | 0.494031 0.123429 |
| 108865.1 | 108865 Zinkstearat Av. of | 0.486698 0.029438 |
| 115419.1 | 115419 Cetylpalmitat Av. of | 0.230904 0.147221 |
| 118308.1 | 118308 Decan-1-sulfons,,ure Na | 0.504881 0.666383 |
| 500090.1 | 500090 L(+)-Ascorbylpalmitat | 0.429206 0.084882 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 101407.1 | 101407 Kristallviolet C.I. 42 | 0.980783 |

| Overlapping with: | Hit |
|--|---------------------|
| 115940.1 115940 Kristallviolet C.I. 42 | 0.017617 0.308797 |

| File Name | Sample Name | Confidence Region |
|-----------|--------------------------------|-------------------|
| 102070.1 | 102070 Calciumchlorid 90-95% | 1.354308 |

| Overlapping with: | Hit |
|--|---------------------|
| 100492.1 100492 Oxals„ure-Dihydrat rei | 1.321093 0.192883 |
| 100807.1 100807 Trichloressigs„ure A | 1.308760 0.217889 |
| 100963.1 100963 DL-Alanin f•r bioch. Z | 1.351249 0.047640 |
| 101190.1 101190 Di-Ammoniumoxalat-Mono | 1.354168 0.222898 |
| 101226.1 101226 Ammoniummonovanadat z. | 1.352215 0.034159 |
| 102146.1 102146 Calciumhydrogenphospha | 1.350569 0.221331 |
| 102148.1 102148 Calcium-D-Saccharat re | 1.341081 0.138641 |
| 102851.1 102851 Calciumlaevulinat rein | 1.329423 0.096867 |
| 104128.1 104128 Auxillin (Gibberillin | 1.311841 0.142809 |
| 104350.1 104350 L-Histidinmonohydrochl | 1.177595 0.415788 |
| 106467.1 106467 Natriumhydroxi granuli | 1.038939 0.417924 |
| 106544.1 106544 Natriumnitrit reinst | 1.295365 0.205559 |
| 107959.1 107959 S•holzextrakt spr•hge | 1.311896 3.519427 |
| 108371.1 108371 L-Tyrosin f. b. Zw. | 1.246063 0.173634 |
| 112428.1 112428 Amygdalin f. Bioch. Zw | 1.292664 0.969447 |
| 159605.1 159605 4-Aminophenol Av. of | 1.265652 0.043963 |
| 271436.1 271436 Adenosin-5-diphosphors | 1.351799 0.178478 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 102072.1 | 102072 Calciumchlorid-Hexahyd | 1.431255 |

| Overlapping with: | Hit |
|--|---------------------|
| 100010.1 100010 Acetanilid reinst kris | 1.288162 0.082312 |
| 100085.1 100085 Acetylsalicyls„ure A | 1.320809 0.055139 |
| 100090.1 100090 Adipins„ure reinst A | 1.402201 0.044166 |
| 100102.1 100102 4-Aminobenzues„ure rei | 1.425541 0.070165 |

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| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 102127.1 | 102127 Beta-Cyclodextrin Av | 0.807182 |

| Overlapping with: | Hit |
|--|---------------------|
| 103006.1 103006 Dextrin wei• Av. of | 0.514819 0.147478 |
| 111685.1 111685 St„rke (aus Weizen) | 0.516588 0.168958 |
| 501606.1 501606 D+Biotin-konz. 2% gesp | 0.598752 0.194068 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 102530.1 | 102530 Cobalt-II-acetat Tetra | 1.291127 |

| Overlapping with: | Hit |
|--|---------------------|
| 103962.1 103962 Eisen(II)-fumarat, rei | 0.311091 0.336673 |
| 107959.1 107959 S•holzextrakt spr•hge | 1.249914 3.519427 |

| File Name | Sample Name | Confidence Region |
|-----------|--------------------------------|-------------------|
| 103006.1 | 103006 Dextrin wei• Av. of | 0.147478 |

| Overlapping with: | Hit |
|---|---------------------|
| 111685.1 111685 St„rke (aus Weizen) | 0.130752 0.168958 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 103962.1 | 103962 Eisen(II)-fumarat, rei | 0.336673 |

| Overlapping with: | Hit |
|--|---------------------|
| 102530.1 102530 Cobalt-II-acetat Tetra | 0.311091 1.291127 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 104168.1 | 104168 Beta-Glycerophosphat D | 1.094834 |

| Overlapping with: | Hit |
|-------------------|-----|
|-------------------|-----|

| | | | | |
|----------|--------|-------------------------|----------|----------|
| 102146.1 | 102146 | Calciumhydrogenphospha | 0.880818 | 0.221331 |
| 110550.1 | 110550 | Creatinphosphors,,ure D | 1.047801 | 0.716645 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 104349.1 | 104349 Homatropinmethylbromid | 1.304249 |

| Overlapping with: | | Hit | |
|-------------------|----------------------------------|----------|----------|
| 100010.1 | 100010 Acetanilid reinst kris | 1.144573 | 0.082312 |
| 100102.1 | 100102 4-Aminobenzues,,ure rei | 1.296015 | 0.070165 |
| 100126.1 | 100129 L-Asparagins,,ure reins | 1.219044 | 0.092033 |
| 100130.1 | 100130 Benzoes,,ure gepulvert | 1.090488 | 0.055978 |
| 100289.1 | 100289 L-Glutamin f. bioch. Z | 1.217203 | 0.176502 |
| 101155.1 | 101155 Di-Ammoniumhydrogencit | 1.300302 | 0.818903 |
| 101838.1 | 101838 Curco BS Av. of 6 | 1.237595 | 0.155954 |
| 102166.1 | 102166 D(-)-Campher raffinier | 1.279589 | 0.073816 |
| 102424.1 | 102424 Chloramin T Trihydrat | 1.252674 | 0.394545 |
| 102425.1 | 102425 Chloralhydrat Av. of | 1.265094 | 0.084440 |
| 102584.1 | 102584 Coffein rein Av. of | 1.207450 | 0.080836 |
| 102838.1 | 102838 L-Cystein f•r bioch. Z | 1.199320 | 0.241116 |
| 103149.1 | 103149 (+-)-Ephedrinhydrochlo | 1.109581 | 0.090064 |
| 104343.1 | 104343 Methenamin Av. of 8 | 1.101072 | 0.097719 |
| 104506.1 | 104506 L-Hydroxyprolin f. b. | 1.036860 | 3.085978 |
| 105995.1 | 105995 (-)-Menthol krist Av | 1.302672 | 0.949150 |
| 106129.1 | 106129 3-Morpholinopropansulf | 1.209550 | 0.161772 |
| 107069.1 | 107069 Phloroglucin (1,3,5-Tr | 1.235970 | 0.281414 |
| 107230.1 | 107230 Phenolphthalein Av. | 1.265307 | 0.102357 |
| 107256.1 | 107256 L-Penylalanin f. b. Zw | 1.075289 | 0.084684 |
| 107260.1 | 107260 Phenylsalicylat Av. | 1.255347 | 0.107905 |
| 107590.1 | 107590 Resorcin reinst Av. | 1.272124 | 0.114703 |
| 107606.1 | 107606 2-desoxy-D-Ribose Av | 1.183087 | 0.089816 |
| 107769.1 | 107769 L-Serin Av. of 8 | 1.204478 | 0.162600 |
| 110220.1 | 110220 Piperazin-1,4-bis(etha | 1.295696 | 0.691588 |
| 111037.1 | 111037 Trinatriumcitrat Av. | 1.271296 | 0.158675 |
| 112428.1 | 112428 Amygdalin f. Bioch. Zw | 1.095353 | 0.969447 |
| 115230.1 | 115230 3-[4-(2-Hydroxyethyl)- | 1.187180 | 0.157989 |
| 907327.1 | 907327 Piperazin-Hexahydrat | 1.272478 | 0.703618 |

| File Name | Sample Name | Confidence Region |
|-----------|--------------------------------|-------------------|
| 104506.1 | 104506 L-Hydroxyprolin f. b. | 3.085978 |

| Overlapping with: | | Hit | |
|-------------------|----------------------------------|----------|----------|
| 100010.1 | 100010 Acetanilid reinst kris | 1.435668 | 0.082312 |
| 100085.1 | 100085 Acetylsalicyls,,ure A | 1.467487 | 0.055139 |
| 100090.1 | 100090 Adipins,,ure reinst A | 1.491228 | 0.044166 |
| 100102.1 | 100102 4-Aminobenzues,,ure rei | 1.365937 | 0.070165 |

262

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105072.1 | 105072 Di-Kaliumoxalat-Monohy | 3.055652 |

| Overlapping with: | | Hit | |
|-------------------|----------------------------------|----------|----------|
| 100010.1 | 100010 Acetanilid reinst kris | 1.325698 | 0.082312 |
| 100085.1 | 100085 Acetylsalicyls,,ure A | 1.354947 | 0.055139 |
| 100090.1 | 100090 Adipins,,ure reinst A | 1.391741 | 0.044166 |
| 100102.1 | 100102 4-Aminobenzues,,ure rei | 1.372636 | 0.070165 |

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| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105102.1 | 105102 Tri-Kaliumphosphat-Tri | 0.683779 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 101124.1 | 101124 Ammoniumdihydrogenphos | 0.593942 | 0.207669 |

| File Name | Sample Name | Confidence Region |
|-----------|-------------|-------------------|
|-----------|-------------|-------------------|

| | | | |
|----------|--------|------------------------|----------|
| 105700.1 | 105700 | L-Lysinmonohydrochlori | 1.397607 |
|----------|--------|------------------------|----------|

| Overlapping with: | | Hit | |
|-------------------|--------|------------------------|-------------------|
| 100103.1 | 100103 | Amidoschwefels„ure z.A | 1.322880 0.046911 |
| 100126.1 | 100129 | L-Asparagins„ure reins | 1.348678 0.092033 |
| 100138.1 | 100138 | 3,5-Dinitrobenzoes„ure | 1.393947 0.032759 |
| 100242.1 | 100242 | Citronens„ure Monohydr | 1.368592 0.458916 |

87

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105870.1 | 105870 Magnesiumhydroxid, rei | 3.057661 |

| Overlapping with: | | Hit | |
|-------------------|--------|------------------------|-------------------|
| 100010.1 | 100010 | Acetanilid reinst kris | 1.411926 0.082312 |
| 100085.1 | 100085 | Acetylsalicyls„ure A | 1.411502 0.055139 |
| 100090.1 | 100090 | Adipins„ure reinst A | 1.414443 0.044166 |
| 100102.1 | 100102 | 4-Aminobenzues„ure rei | 1.412764 0.070165 |

262

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105876.1 | 105876 Magnesiumstearat gef„l | 0.123470 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 101107.1 | 101107 Aluminiumstearat Av. | 0.103617 | 0.075229 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105924.1 | 105924 Mangan-II-carbonat-Hyd | 0.761684 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 102071.1 | 102071 Calciumcarbonat f. op. | 0.661065 | 0.011004 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 105995.1 | 105995 (-)-Menthol krist Av | 0.949150 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 500859.1 | 500859 Vitamin E-Trockenpulve | 0.934005 | 0.094467 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 106432.1 | 106432 Tri-Natriumcitrat-Dihy | 0.178043 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 106446.1 | 106446 tri-Natriumcitrat Dihy | 0.036992 | 0.249648 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 106446.1 | 106446 tri-Natriumcitrat Dihy | 0.249648 |

| Overlapping with: | | Hit | |
|-------------------|---------------------------------|----------|----------|
| 106432.1 | 106432 Tri-Natriumcitrat-Dihy | 0.036992 | 0.178043 |

| File Name | Sample Name | Confidence Region |
|-----------|---------------------------------|-------------------|
| 107428.1 | 107428 Propyl-4-hydroxybenzoa | 1.665151 |

| Overlapping with: | | Hit | |
|-------------------|--------|------------------------|-------------------|
| 100010.1 | 100010 | Acetanilid reinst kris | 1.395932 0.082312 |
| 100085.1 | 100085 | Acetylsalicyls„ure A | 1.464164 0.055139 |
| 100090.1 | 100090 | Adipins„ure reinst A | 1.423003 0.044166 |
| 100102.1 | 100102 | 4-Aminobenzues„ure rei | 1.295012 0.070165 |

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| File Name | Sample Name | Confidence | Region |
|-------------------|---------------------------------|------------|----------|
| 107656.1 | 107656 Lactose-Monohydrat gep | 0.067694 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 501551.1 | 501551 Vitamin D3-Trockenpulv | 0.036849 | 0.081402 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 107959.1 | 107959 S-áholzextrakt spr•hge | 3.519427 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 100010.1 | 100010 Acetanilid reinst kris | 1.414047 | 0.082312 |
| 100085.1 | 100085 Acetylsalicyls„ure A | 1.394373 | 0.055139 |
| 100090.1 | 100090 Adipins„ure reinst A | 1.414481 | 0.044166 |
| 100102.1 | 100102 4-Aminobenzues„ure rei | 1.418794 | 0.070165 |
| ----- | | | |
| 262 | | | |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 108198.1 | 108198 Tablettierhilsmittel T | 0.559912 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 108196.1 | 108196 Tablettierhilsmittel D | 0.393236 | 0.306419 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 108337.1 | 108337 D+Glucose wasserfrei | 0.069033 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 278343.1 | 278343 Dextroseandydrid Av. | 0.042041 | 0.081600 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 108484.1 | 108484 Harnstoff, Perlform re | 0.131105 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 108486.1 | 108486 Harnstoff reinst krist | 0.015151 | 0.084571 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 108486.1 | 108486 Harnstoff reinst krist | 0.084571 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 108484.1 | 108484 Harnstoff, Perlform re | 0.015151 | 0.131105 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 111352.1 | 111352 Molinon Instant Av. | 0.695643 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 105321.1 | 105321 D-(-)-Fructose reinst | 0.475020 | 0.180837 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 111685.1 | 111685 St„rke (aus Weizen) | 0.168958 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 103006.1 | 103006 Dextrin weiá Av. of | 0.130752 | 0.147478 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 112428.1 | 112428 Amygdalin f. Bioch. Zw | 0.969447 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 100010.1 | 100010 Acetanilid reinst kris | 0.942432 | 0.082312 |
| 100130.1 | 100130 Benzoes„ure gepulvert | 0.956024 | 0.055978 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 115940.1 | 115940 Kristallviolet C.I. 42 | 0.308797 | |
| ----- | | | |
| Overlapping with: | | Hit | |
| 101407.1 | 101407 Kristallviolet C.I. 42 | 0.017617 | 0.980783 |
| ----- | | | |
| File Name | Sample Name | Confidence | Region |
| 118305.1 | 118305 Hexan-1-sulfons„ure Na | 0.739253 | |

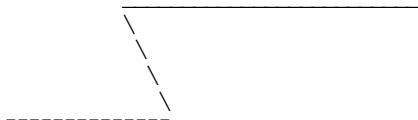
| | | | | |
|-------------------|-------------|-------------------------|------------|----------|
| ----- | | | | |
| Overlapping with: | | | | |
| 118304.1 | 118304 | Pentan-1-Sulfons,,urena | Hit | |
| 118306.1 | 118306 | Heptan-1-sulfons,,ure N | 0.637670 | 0.529805 |
| 118306.1 | 118306 | Heptan-1-sulfons,,ure N | 0.375709 | 0.581004 |
| File Name | Sample Name | | Confidence | Region |
| 118306.1 | 118306 | Heptan-1-sulfons,,ure N | | 0.581004 |
| ----- | | | | |
| Overlapping with: | | | | |
| 118305.1 | 118305 | Hexan-1-sulfons,,ure Na | Hit | |
| 118307.1 | 118307 | Octan-1-sulfons,,ure Na | 0.375709 | 0.739253 |
| 118307.1 | 118307 | Octan-1-sulfons,,ure Na | 0.540796 | 0.804557 |
| File Name | Sample Name | | Confidence | Region |
| 118307.1 | 118307 | Octan-1-sulfons,,ure Na | | 0.804557 |
| ----- | | | | |
| Overlapping with: | | | | |
| 100671.1 | 100671 | Stearins,,ure Av. of | Hit | |
| 100989.1 | 100989 | Cetylalkohol (1-Hexade | 0.670867 | 0.026060 |
| 101107.1 | 101107 | Aluminiumstearat Av. | 0.717941 | 0.589218 |
| 102154.1 | 102154 | Calciumstearat Av. o | 0.546261 | 0.075229 |
| 105876.1 | 105876 | Magnesiumstearat gef,,l | 0.716758 | 0.107628 |
| 107682.1 | 107682 | Seife medizinisch Av | 0.534584 | 0.123470 |
| 108865.1 | 108865 | Zinkstearat Av. of | 0.547598 | 0.123429 |
| 115419.1 | 115419 | Cetylpalmitat Av. of | 0.739485 | 0.029438 |
| 118305.1 | 118305 | Hexan-1-sulfons,,ure Na | 0.626674 | 0.147221 |
| 118306.1 | 118306 | Heptan-1-sulfons,,ure N | 0.778760 | 0.739253 |
| 118308.1 | 118308 | Decan-1-sulfons,,ure Na | 0.540796 | 0.581004 |
| 500090.1 | 500090 | L(+)-Ascorbylpalmitat | 0.402931 | 0.666383 |
| 500090.1 | 500090 | L(+)-Ascorbylpalmitat | 0.615216 | 0.084882 |
| File Name | Sample Name | | Confidence | Region |
| 118308.1 | 118308 | Decan-1-sulfons,,ure Na | | 0.666383 |
| ----- | | | | |
| Overlapping with: | | | | |
| 100671.1 | 100671 | Stearins,,ure Av. of | Hit | |
| 100989.1 | 100989 | Cetylalkohol (1-Hexade | 0.396762 | 0.026060 |
| 101107.1 | 101107 | Aluminiumstearat Av. | 0.504881 | 0.589218 |
| 102154.1 | 102154 | Calciumstearat Av. o | 0.261219 | 0.075229 |
| 105876.1 | 105876 | Magnesiumstearat gef,,l | 0.439122 | 0.107628 |
| 107682.1 | 107682 | Seife medizinisch Av | 0.248754 | 0.123470 |
| 108865.1 | 108865 | Zinkstearat Av. of | 0.460189 | 0.123429 |
| 115419.1 | 115419 | Cetylpalmitat Av. of | 0.530342 | 0.029438 |
| 118307.1 | 118307 | Octan-1-sulfons,,ure Na | 0.385547 | 0.147221 |
| 500090.1 | 500090 | L(+)-Ascorbylpalmitat | 0.402931 | 0.804557 |
| 500090.1 | 500090 | L(+)-Ascorbylpalmitat | 0.421376 | 0.084882 |
| File Name | Sample Name | | Confidence | Region |
| 278343.1 | 278343 | Dextroseandhydrid Av. | | 0.081600 |
| ----- | | | | |
| Overlapping with: | | | | |
| 108337.1 | 108337 | D+Glucose wasserfrei | Hit | |
| 108337.1 | 108337 | D+Glucose wasserfrei | 0.042041 | 0.069033 |
| File Name | Sample Name | | Confidence | Region |
| 450890.1 | 450890 | Saponin Av. of 10 | | 0.131734 |
| ----- | | | | |
| Overlapping with: | | | | |
| 501551.1 | 501551 | Vitamin D3-Trockenpulv | Hit | |
| 501551.1 | 501551 | Vitamin D3-Trockenpulv | 0.117565 | 0.081402 |
| File Name | Sample Name | | Confidence | Region |
| 451318.1 | 451318 | DL-Methionin Av. of | | 0.987647 |
| ----- | | | | |
| Overlapping with: | | | | |
| 105707.1 | 105707 | L-Methionin Av. of | Hit | |
| 105707.1 | 105707 | L-Methionin Av. of | 0.584818 | 0.034623 |
| File Name | Sample Name | | Confidence | Region |
| 501551.1 | 501551 | Vitamin D3-Trockenpulv | | 0.081402 |
| ----- | | | | |
| Overlapping with: | | | | |
| 107656.1 | 107656 | Lactose-Monohydrat gep | Hit | |
| 107656.1 | 107656 | Lactose-Monohydrat gep | 0.036849 | 0.067694 |
| File Name | Sample Name | | Confidence | Region |
| 102162.1 | 102162 | Calciumsulfat Hemihydr | | 1.337265 |
| ----- | | | | |
| Overlapping with: | | | | |
| 101841.1 | 101841 | Oxals,,ure-bis-(Cyclohe | Hit | |
| 104005.1 | 104005 | Paraformaldehyd reinst | 1.129260 | 0.058678 |
| 104005.1 | 104005 | Paraformaldehyd reinst | 1.302986 | 0.051862 |

| | | | | |
|----------|--------|------------------------|----------|----------|
| 107311.1 | 107311 | Pilocarpinhydrochlorid | 1.287550 | 0.092759 |
| 501902.1 | 501902 | Trihydroxyethylrutin | 1.336778 | 0.299547 |
| 524950.1 | 524950 | Vitamin B12 Av. of 1 | 1.045725 | 0.135492 |

Summary:

Average Spectra, which can be confused with other references: 42 of 263
Average Spectra, which can be uniquely identified: 221 of 263

| | | |
|----------|--------|-------------------------|
| 100010.1 | 100010 | Acetanilid reinst kris |
| 100085.1 | 100085 | Acetylsalicyls„ure A |
| 100090.1 | 100090 | Adipins„ure reinst A |
| 100102.1 | 100102 | 4-Aminobenzoes„ure rei |
| 100103.1 | 100103 | Amidoschwefels„ure z.A |
| 100126.1 | 100129 | L-Asparagins„ure reins |
| 100130.1 | 100130 | Benzoets„ure gepulvert |
| 100138.1 | 100138 | 3,5-Dinitrobenzoets„ure |
| 100162.1 | 100162 | Bors„ure gepulvert rei |
| 100163.1 | 100163 | Di-Bortrioxid Av. of |
| 100241.1 | 100241 | Citronens„ure wasserfr |
| 100289.1 | 100289 | L-Glutamin f. bioch. Z |
| 100290.1 | 100290 | L-Glutamins„ure reinst |



KL2 Bin

| | | |
|-------------------------|--------------------------|--------------------------|
| 12 100242# 13 100243 3 | 78 102530# 155 106688 4 | 149 106446# 147 106432 3 |
| 13 100243# 12 100242 3 | 85 103006# 211 111685 3 | 150 106467# 127 105691 4 |
| 21 100671# 37 101107 2 | 91 103962# 73 102196 3 | 155 106688# 78 102530 4 |
| 21 100671# 132 105876 3 | 91 103962# 114 104982 4 | 155 106688# 91 103962 4 |
| 21 100671# 223 115419 3 | 91 103962# 121 105125 4 | 155 106688# 115 105061 3 |
| 21 100671# 229 118308 4 | 91 103962# 145 106392 4 | 155 106688# 121 105125 4 |
| 33 100989# 223 115419 4 | 91 103962# 155 106688 4 | 174 107656# 237 450890 2 |
| 37 101107# 21 100671 2 | 114 104982# 78 102530 4 | 174 107656# 258 501551 3 |
| 37 101107# 70 102154 3 | 114 104982# 91 103962 4 | 188 108337# 236 278343 4 |
| 37 101107# 132 105876 1 | 114 104982# 145 106392 4 | 201 108865# 70 102154 2 |
| 37 101107# 223 115419 1 | 115 105061# 45 101187 4 | 211 111685# 85 103006 3 |
| 45 101187# 115 105061 4 | 115 105061# 155 106688 3 | 223 115419# 21 100671 3 |
| 53 101407# 224 115940 1 | 121 105125# 78 102530 4 | 223 115419# 33 100989 4 |
| 70 102154# 37 101107 3 | 121 105125# 91 103962 4 | 223 115419# 37 101107 1 |
| 70 102154# 132 105876 4 | 121 105125# 155 106688 4 | 223 115419# 70 102154 4 |
| 70 102154# 201 108865 2 | 127 105691# 150 106467 4 | 223 115419# 132 105876 2 |
| 70 102154# 223 115419 4 | 132 105876# 21 100671 3 | 224 115940# 53 101407 1 |
| 73 102196# 78 102530 3 | 132 105876# 37 101107 1 | 229 118308# 21 100671 4 |
| 73 102196# 91 103962 3 | 132 105876# 70 102154 4 | 236 278343# 188 108337 4 |
| 78 102530# 73 102196 3 | 132 105876# 223 115419 2 | 237 450890# 174 107656 2 |
| 78 102530# 114 104982 4 | 145 106392# 78 102530 4 | 258 501551# 174 107656 3 |
| 78 102530# 121 105125 4 | 145 106392# 91 103962 4 | |
| 78 102530# 145 106392 4 | 145 106392# 114 104982 4 | |
| | 147 106432# 149 106446 3 | |

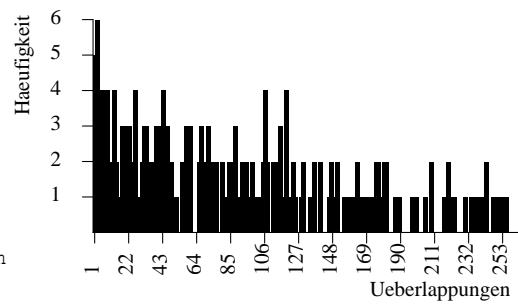
7.4 Matrixdarstellungen

7.4.1 NIR, Standardmethode

```

SV00_MFix.eps
Check Average Spectra
Constant conf. level : Fixed algorithm, Single Sphere Method

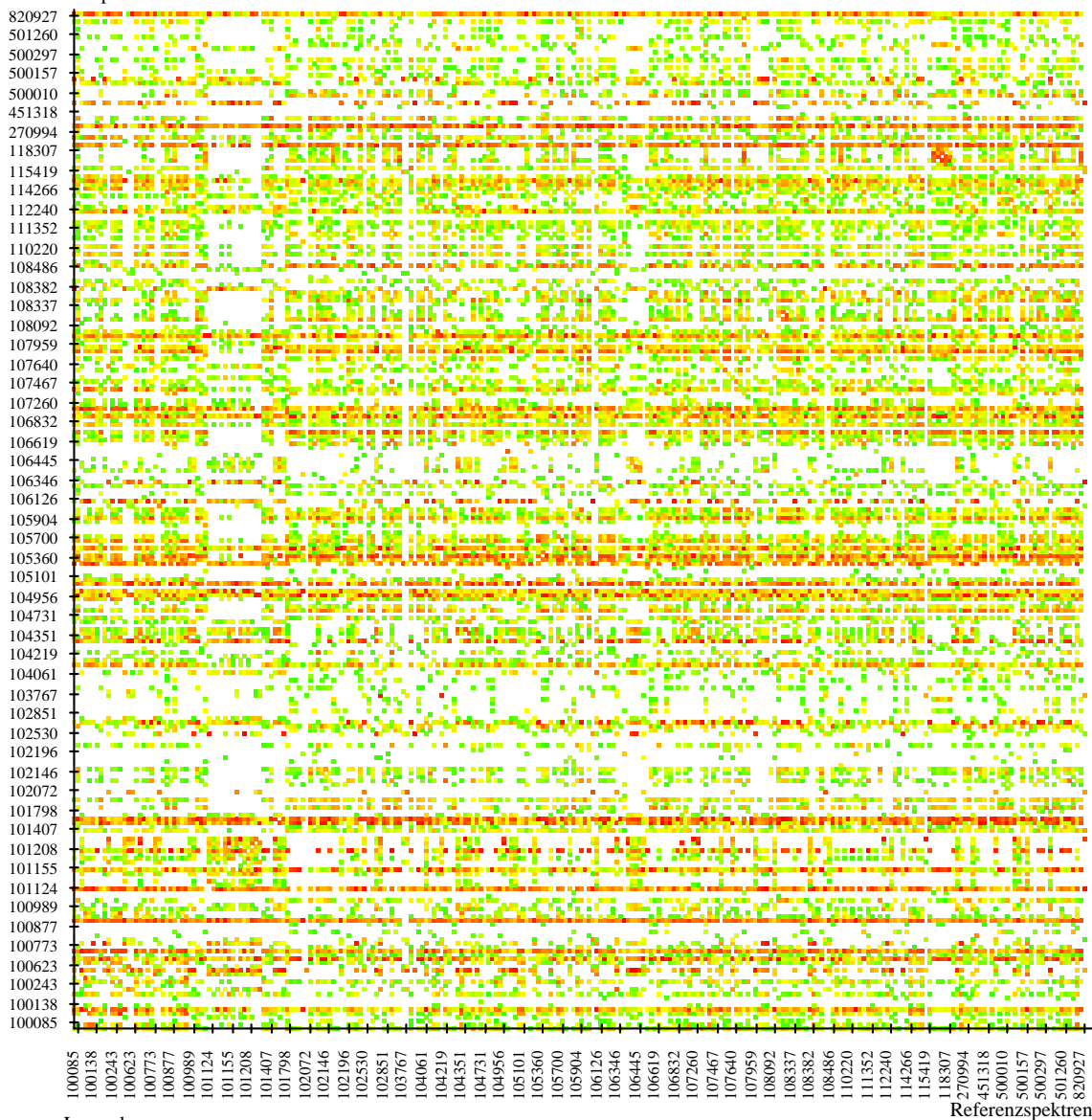
Algorithm:                               Standard
No. of used factor sp.:                   -
Vector normalized spectra:                 Nein
No. of Spectra:                           263
From:                                       3999.71
to:                                         11999.1
Order of Internal Derivation:              0
Smoothing Points for Internal Derivation:  1
    
```



Ueberlapp. Ges.: 23116 (33.55 % bei 1257 (97.7 %) Spektren
 Ueberlapp. Max.:255 bei 1 Spektren

wahr positiv: 45790 falsch negativ: 23116 Sensitivitaet: 66.453 %

Getestete Spektren



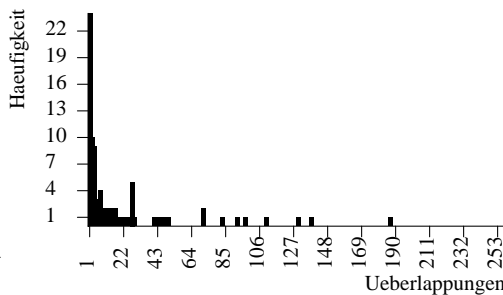
Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N1: Standardmethode ohne Vorbehandlung.

SNV00_MFix.eps
 Check Average Spectra
 Constant conf. level : Fixed algorithm, Single Sphere Method

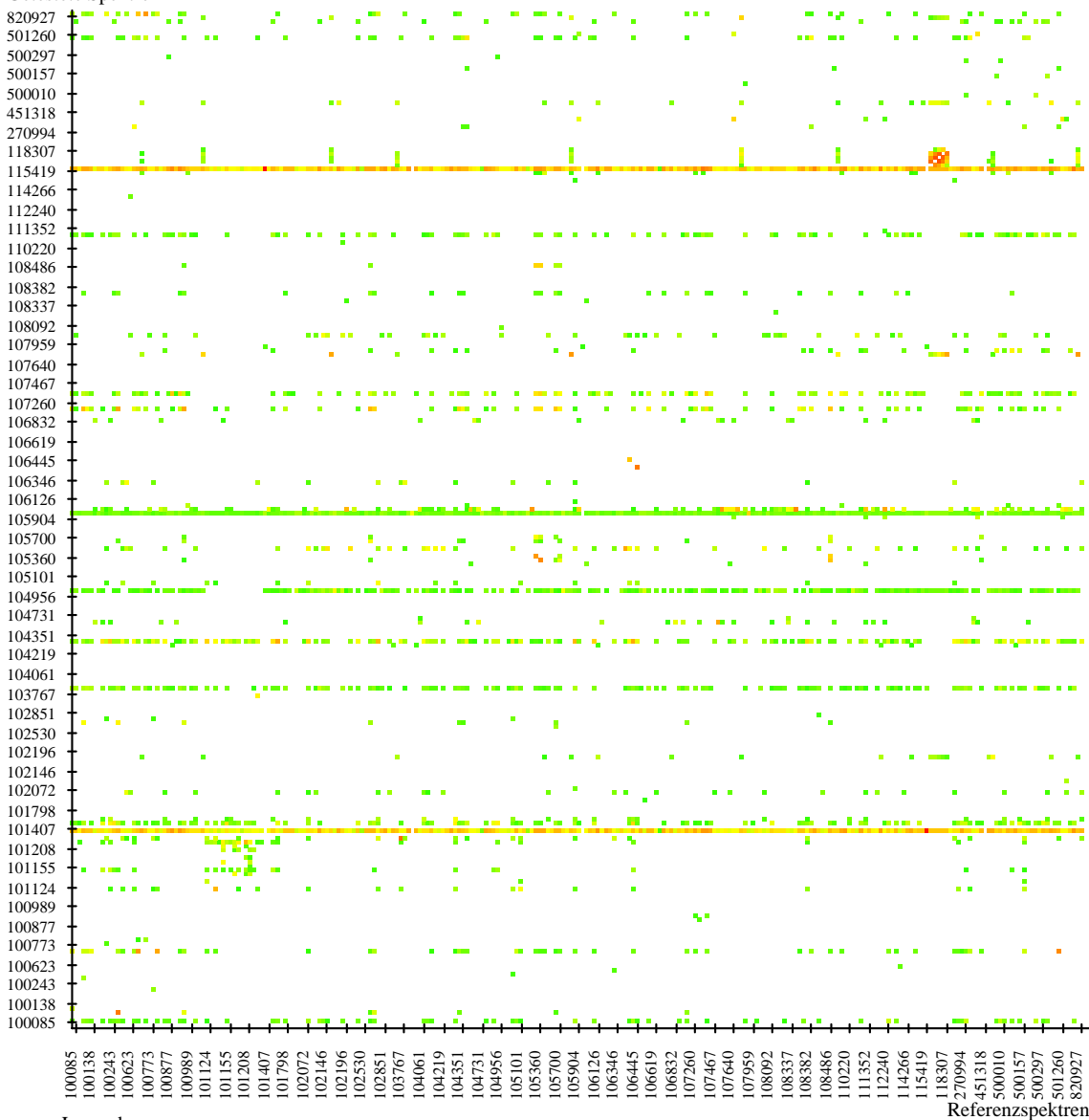
Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 0
 Smoothing Points for Internal Derivation: 1



Ueberlapp. Ges.: 2565 (3.72 %) bei 97 (36.8 %) Spektren
 Ueberlapp. Max.: 260 bei 1 Spektren

wahr positiv: 66341 falsch negativ: 2565 Sensitivitaet: 96.278 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N2: Standardmethode (Vektornormierung).

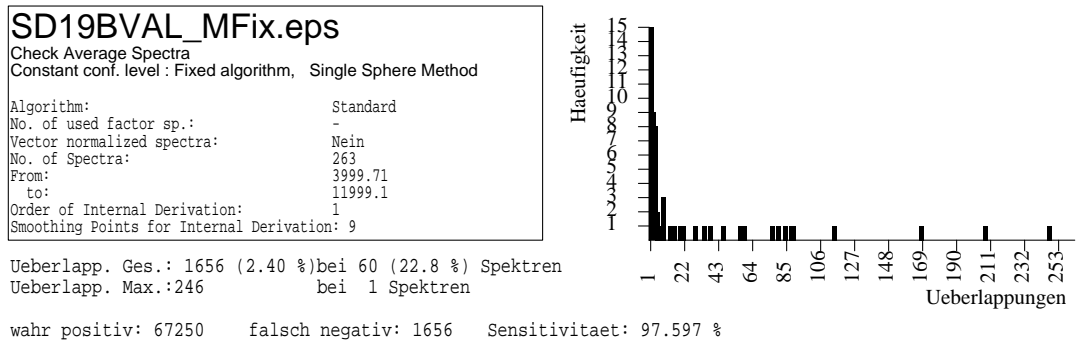


Abb. N3: Standardmethode (1. Ableitung).

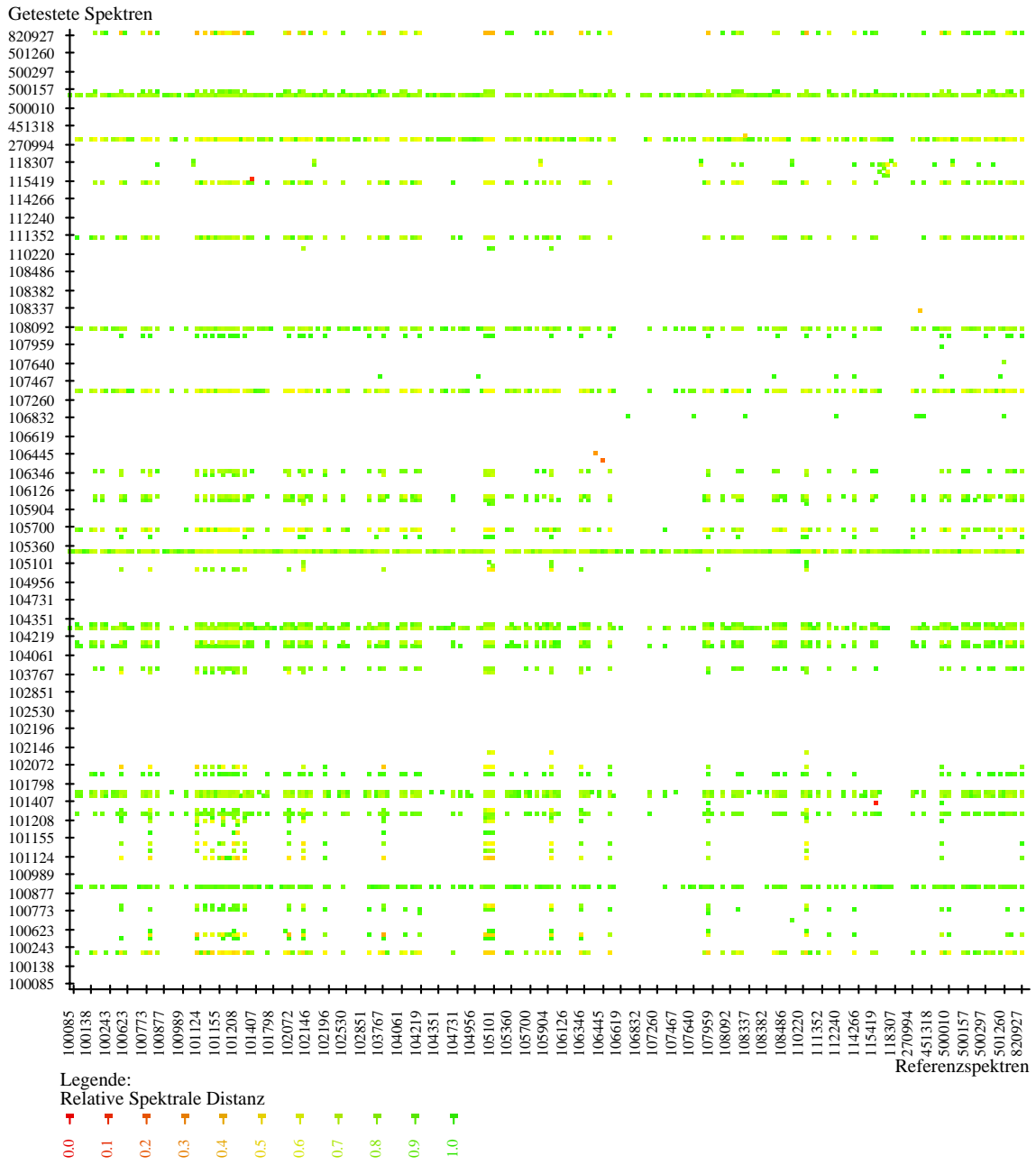
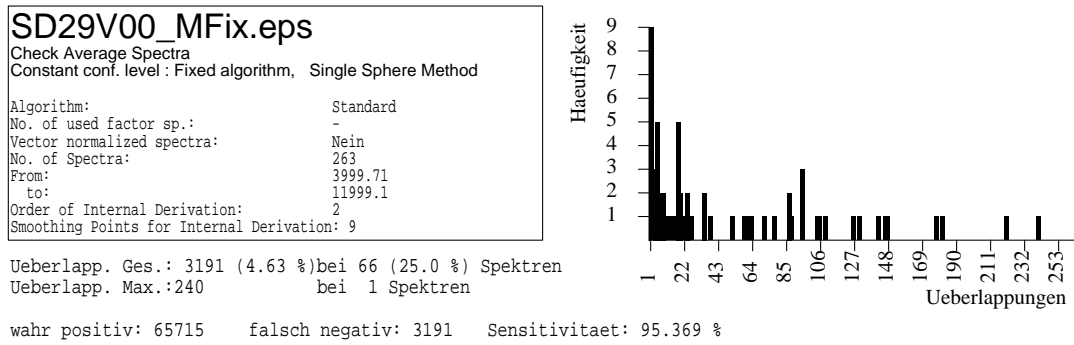
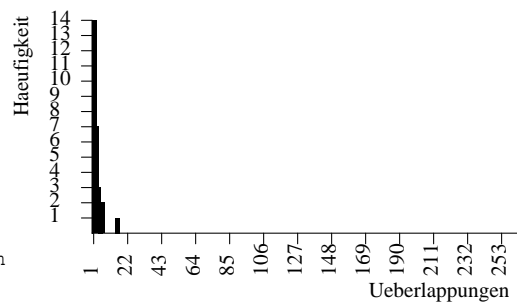


Abb. N4: Standardmethode (2. Ableitung).

SND19dVAL_MFix.eps
 Check Average Spectra
 Constant conf. level : Fixed algorithm, Single Sphere Method

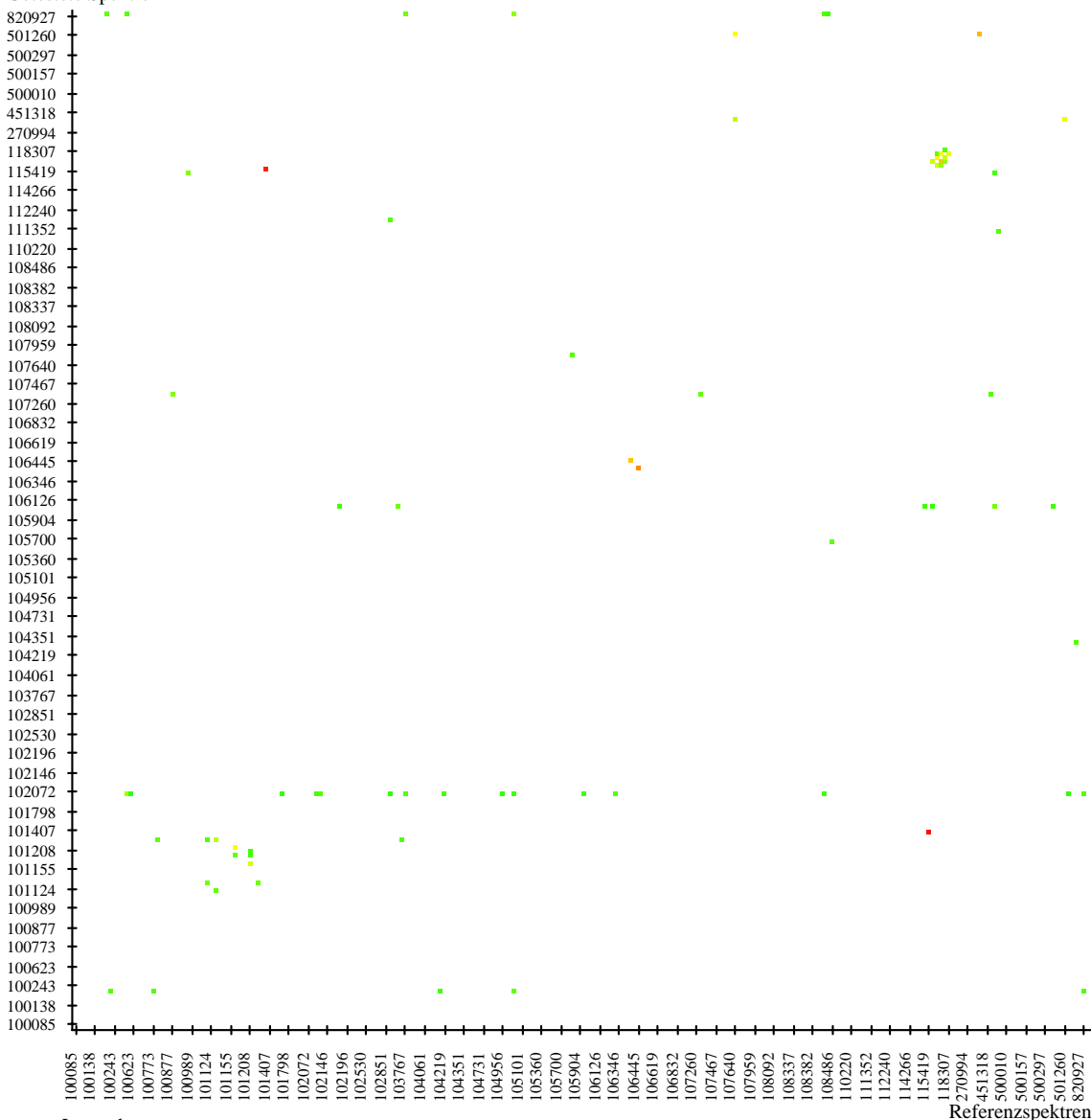
Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 To: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 73 (0.11 %) bei 29 (11.0 %) Spektren
 Ueberlapp. Max.: 15 bei 1 Spektren

wahr positiv: 68833 falsch negativ: 73 Sensitivitaet: 99.894 %

Getestete Spektren



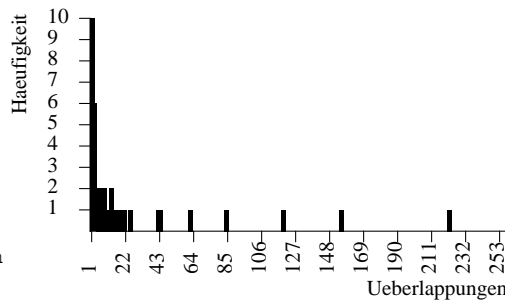
Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N5: Standardmethode (Vektornormierung und 1. Ableitung).

SND29V00_MFix.eps
 Check Average Spectra
 Constant conf. level : Fixed algorithm, Single Sphere Method

Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 2
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 2769 (4.02 %) bei 48 (18.2 %) Spektren
 Ueberlapp. Max.: 260 bei 7 Spektren

wahr positiv: 66137 falsch negativ: 2769 Sensitivitaet: 95.981 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N6: Standardmethode (Vektornormierung und 2. Ableitung).

7.4.2 NIR, Faktormethode

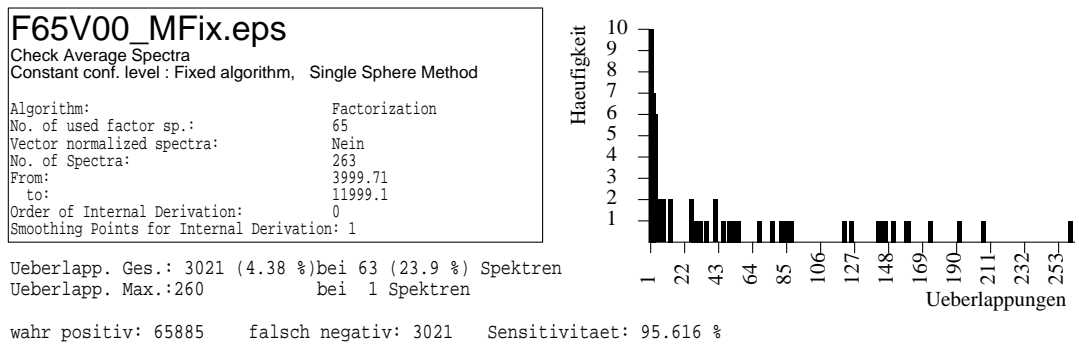
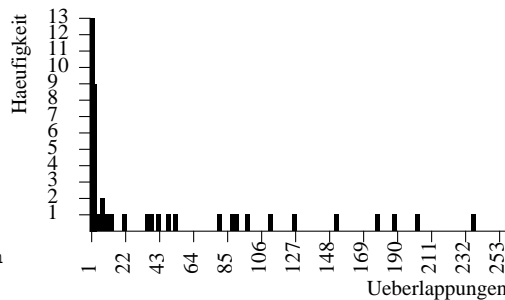


Abb. N7: Faktormethode (65 Faktoren).

F65NV00_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

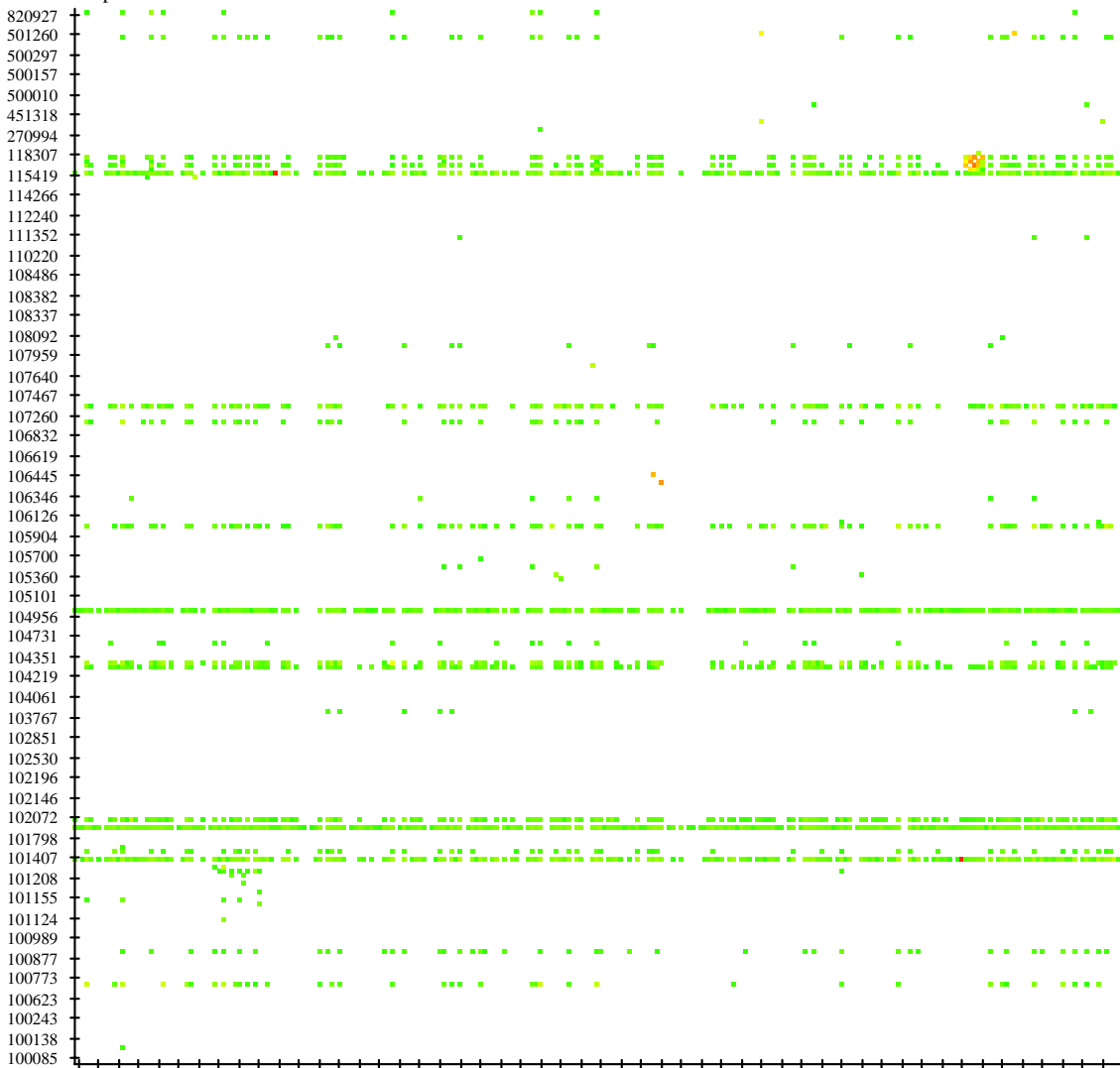
Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 0
 Smoothing Points for Internal Derivation: 1



Ueberlapp. Ges.: 1883 (2.73 %) bei 50 (19.0 %) Spektren
 Ueberlapp. Max.: 236 bei 1 Spektren

wahr positiv: 67023 falsch negativ: 1883 Sensitivitaet: 97.267 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Referenzspektren

Abb. N8: Faktormethode (65 Faktoren, Vektornormierung).

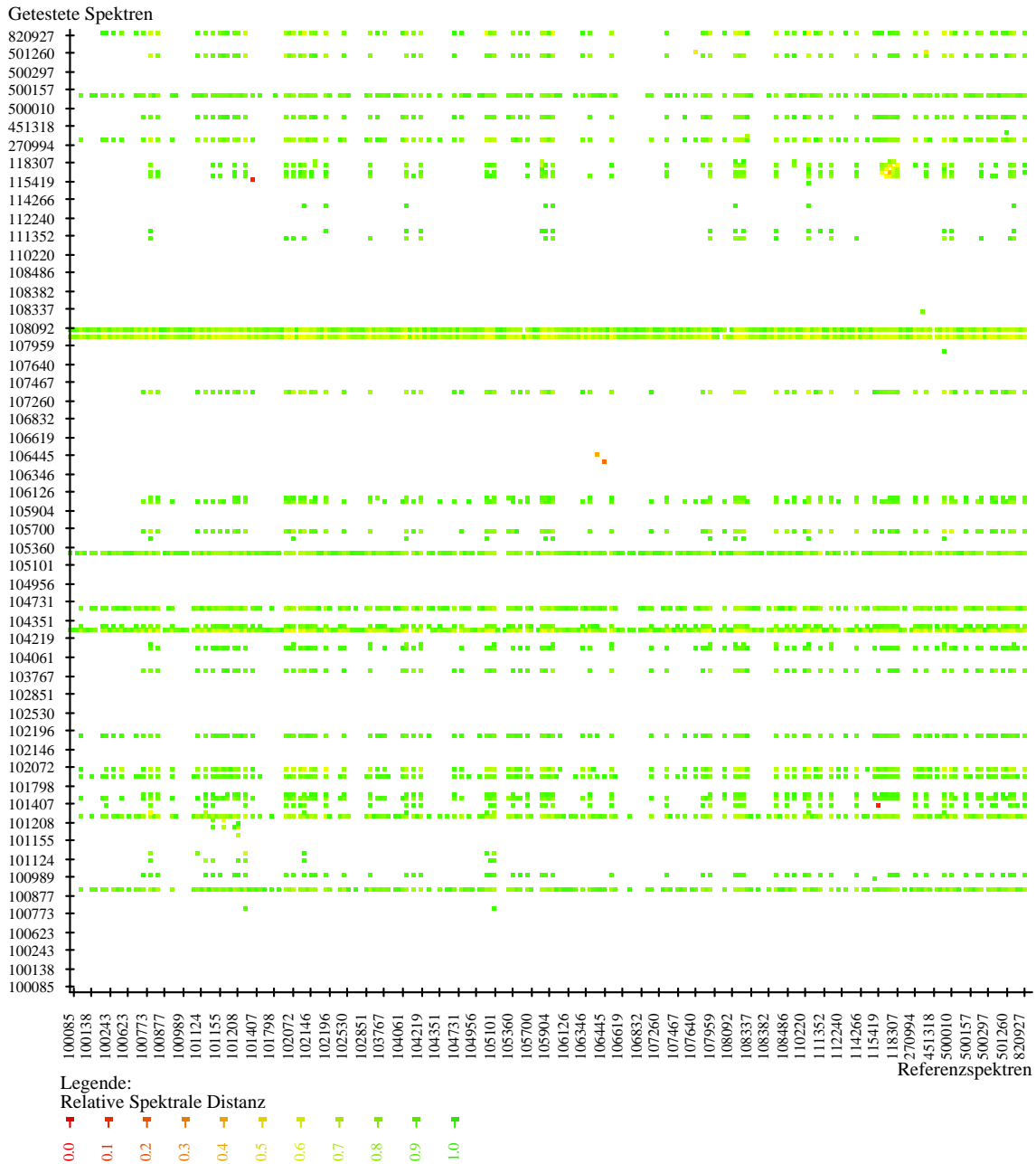
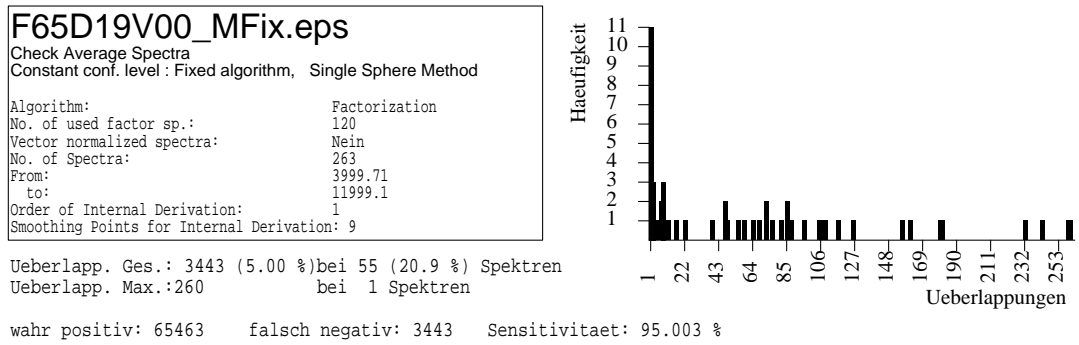


Abb. N9: Faktormethode (65 Faktoren, 1. Ableitung).

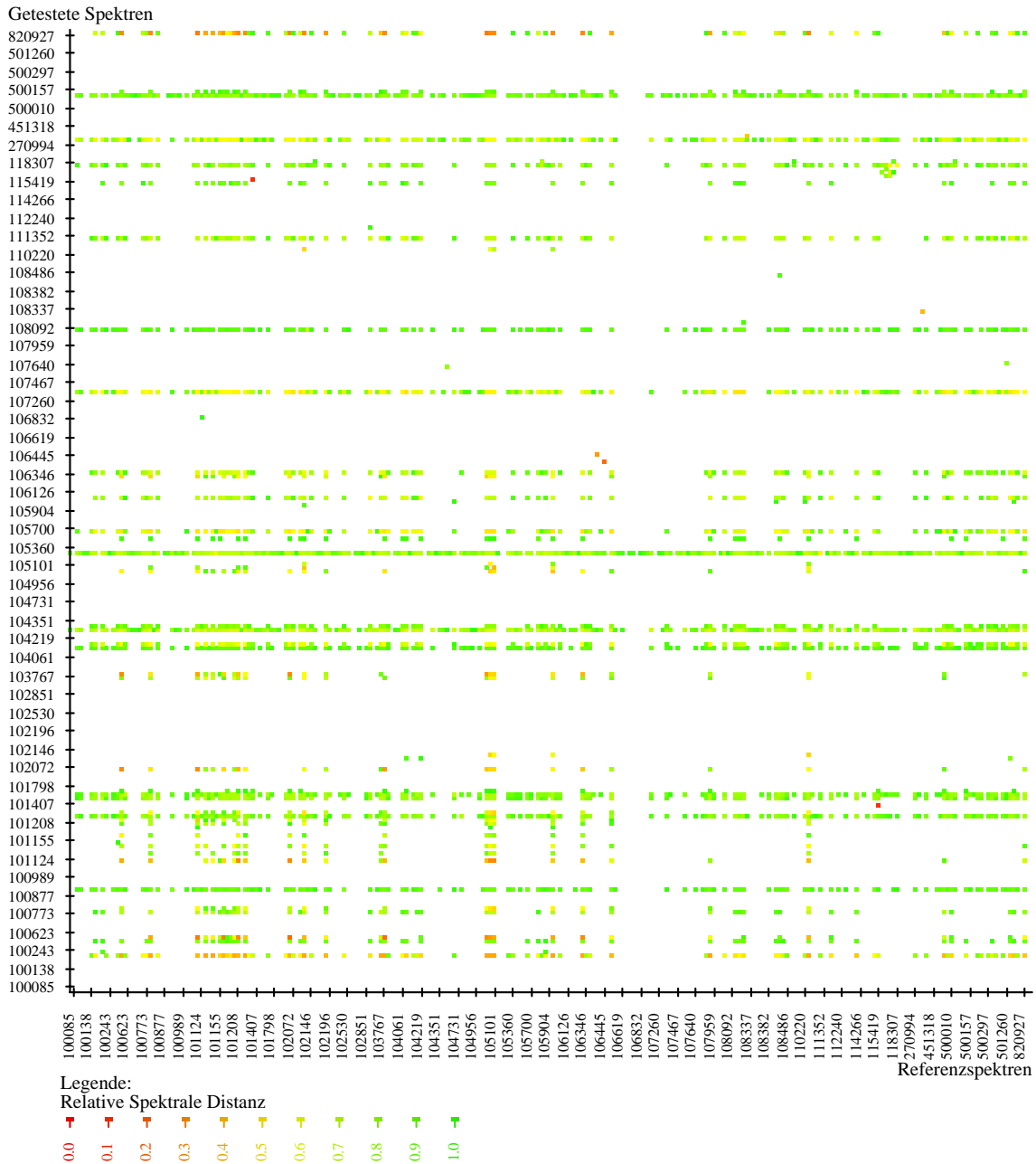
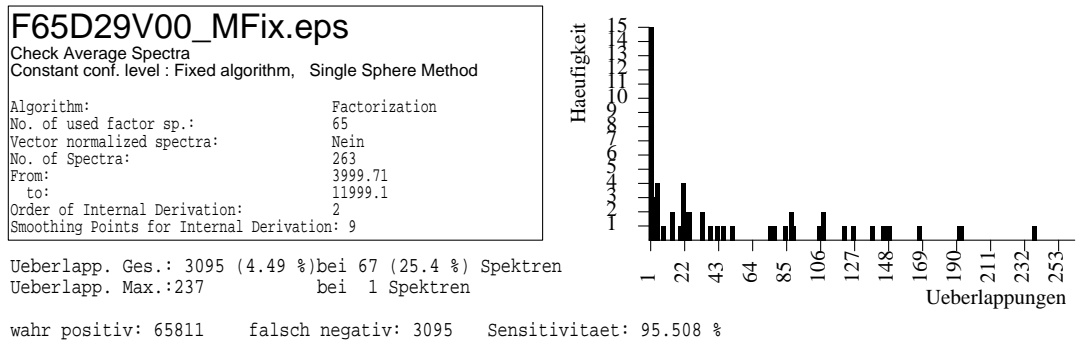
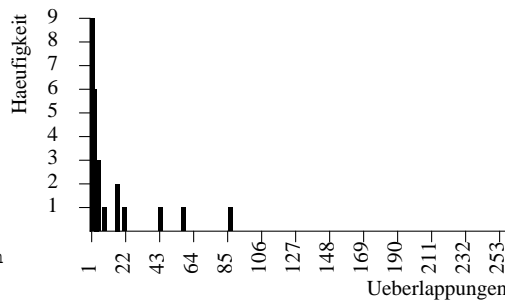


Abb. N10: Faktormethode (65 Faktoren, 2. Ableitung).

F65ND19V00_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 292 (0.42 %) bei 28 (10.6 %) Spektren
 Ueberlapp. Max.: 86 bei 1 Spektren

wahr positiv: 68614 falsch negativ: 292 Sensitivitaet: 99.576 %

Getestete Spektren



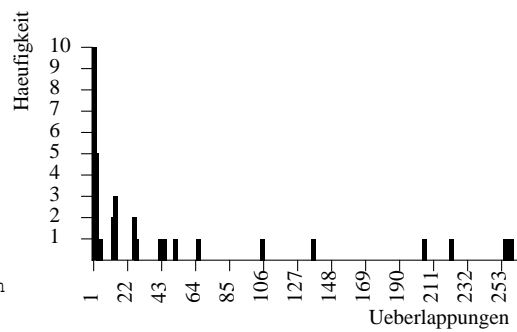
Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N11: Faktormethode (65 Faktoren, Vektornormierung und 1. Ableitung).

f65nd29val_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

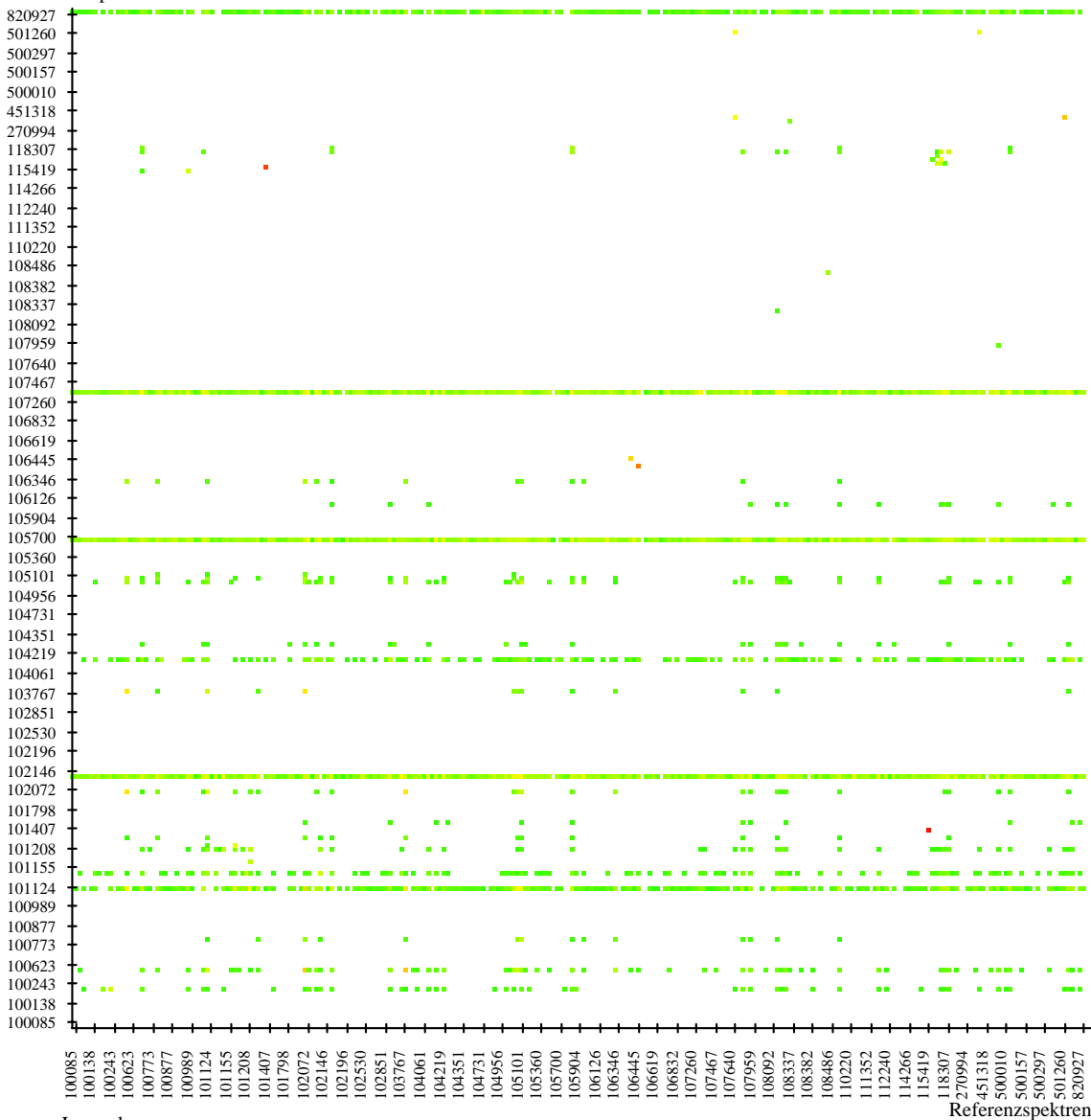
Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 2
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 1841 (2.67 %) bei 39 (14.8 %) Spektren
 Ueberlapp. Max.: 258 bei 1 Spektren

wahr positiv: 67065 falsch negativ: 1841 Sensitivitaet: 97.328 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N12: Faktormethode (65 Faktoren, Vektornormierung und 2. Ableitung).

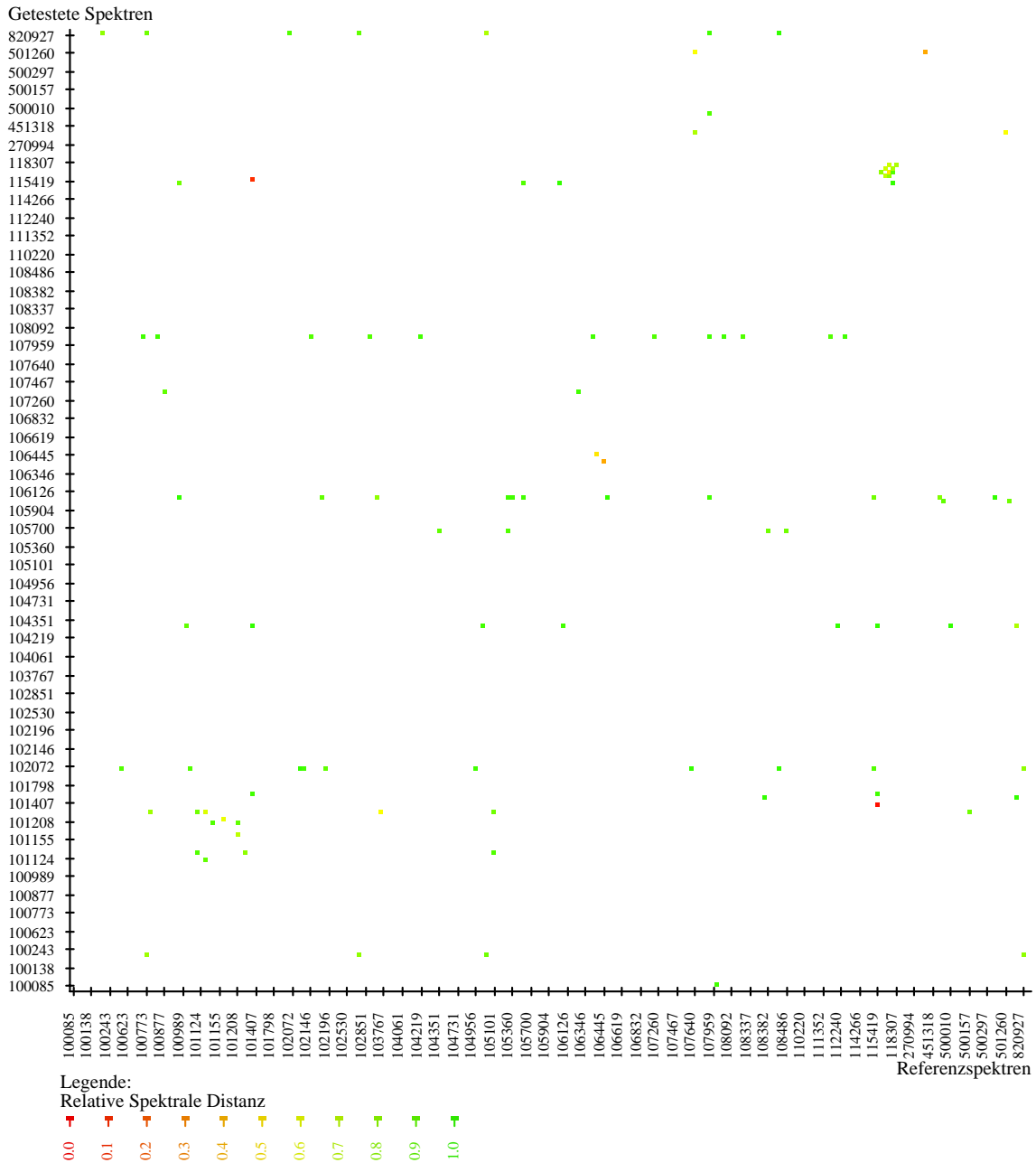
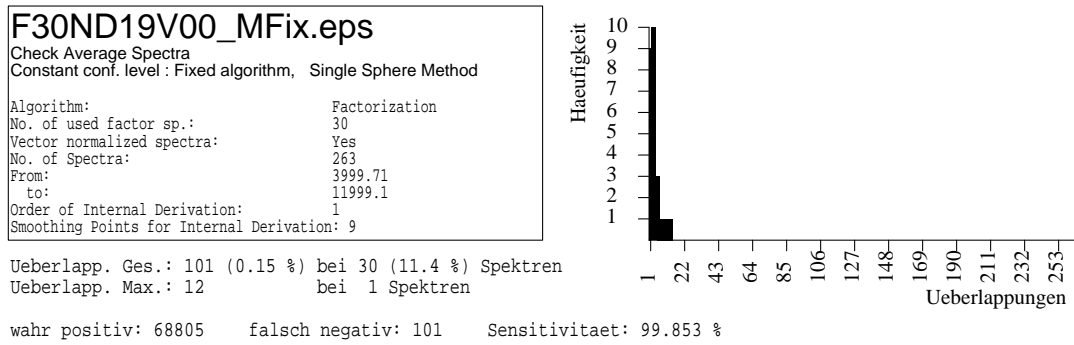


Abb. N13: Faktormethode (30 Faktoren, Vektornormierung und 1. Ableitung).

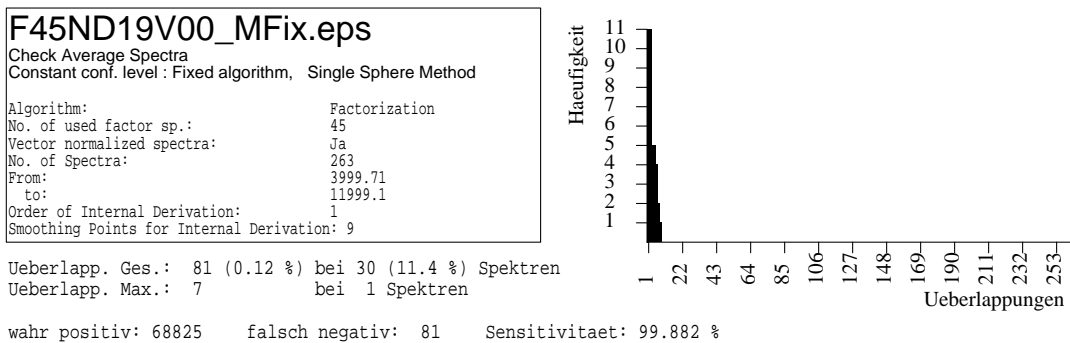


Abb. N14: Faktormethode (45 Faktoren, Vektornormierung und 1. Ableitung).

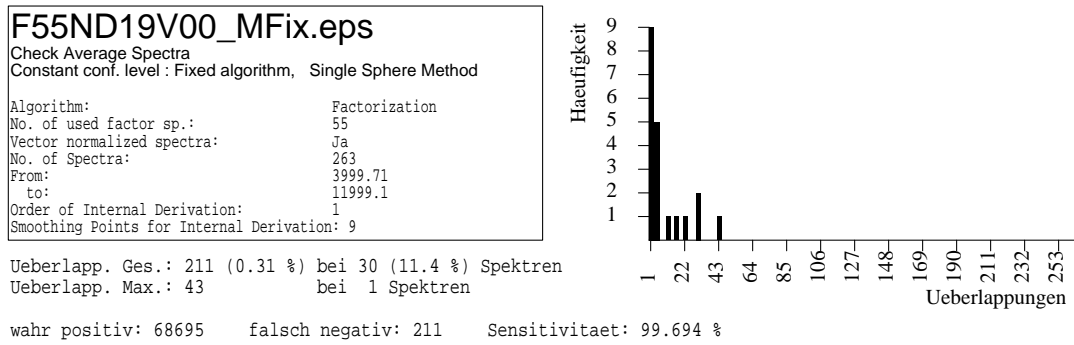
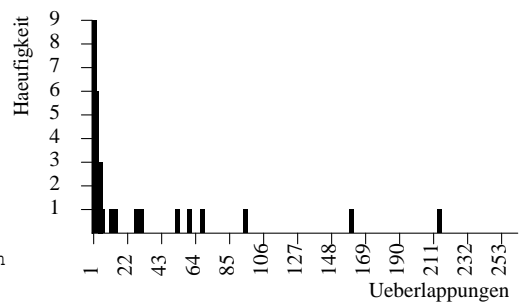


Abb. N15: Faktormethode (55 Faktoren, Vektornormierung und 1. Ableitung).

F90ND19V00_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

Algorithm: Factorization
 No. of used factor sp.: 90
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 782 (1.13 %) bei 32 (12.1 %) Spektren
 Ueberlapp. Max.: 214 bei 1 Spektren

wahr positiv: 68124 falsch negativ: 782 Sensitivitaet: 98.865 %

Getestete Spektren

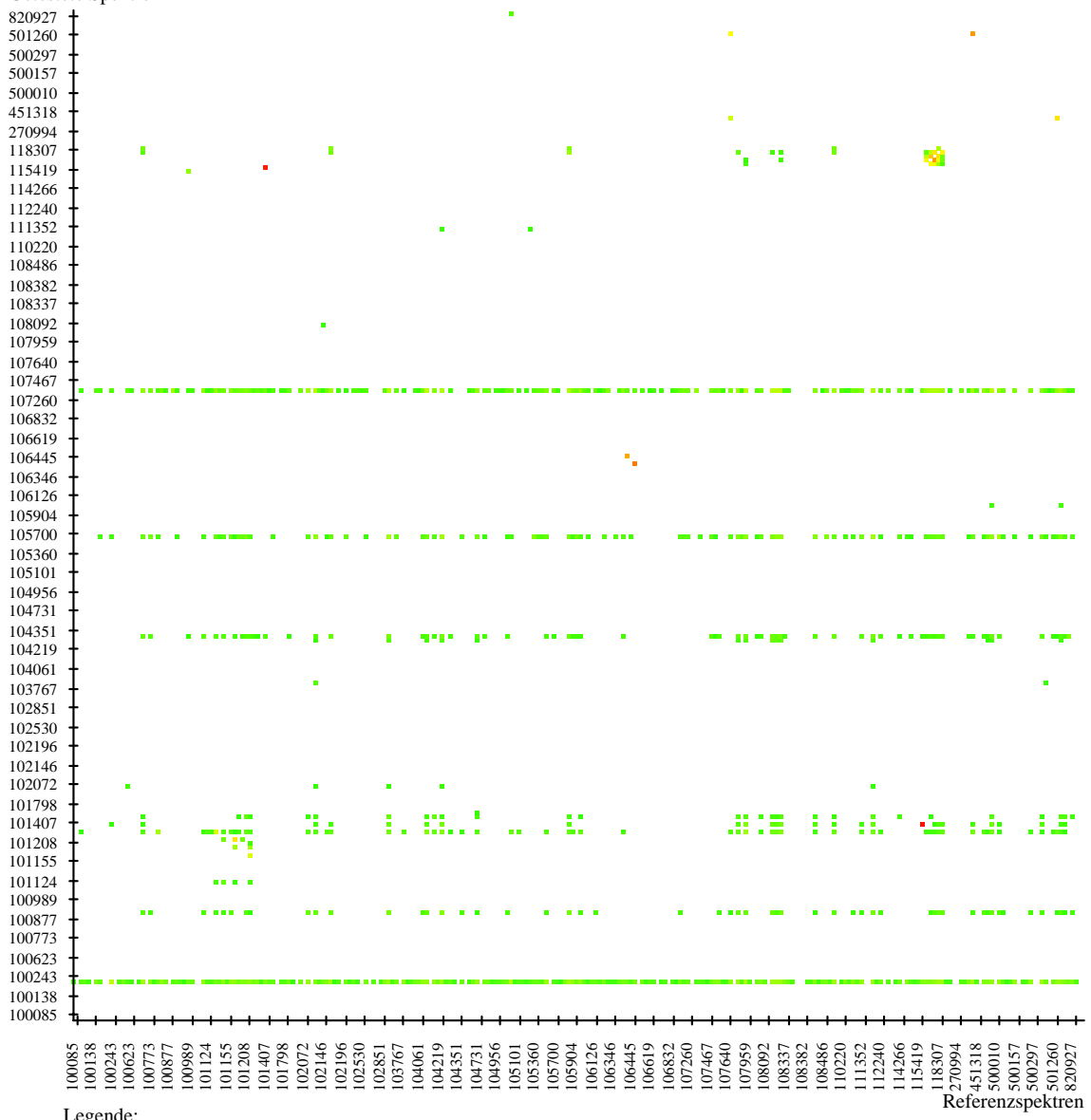


Abb. N16: Faktormethode (90 Faktoren, Vektornormierung und 1. Ableitung).

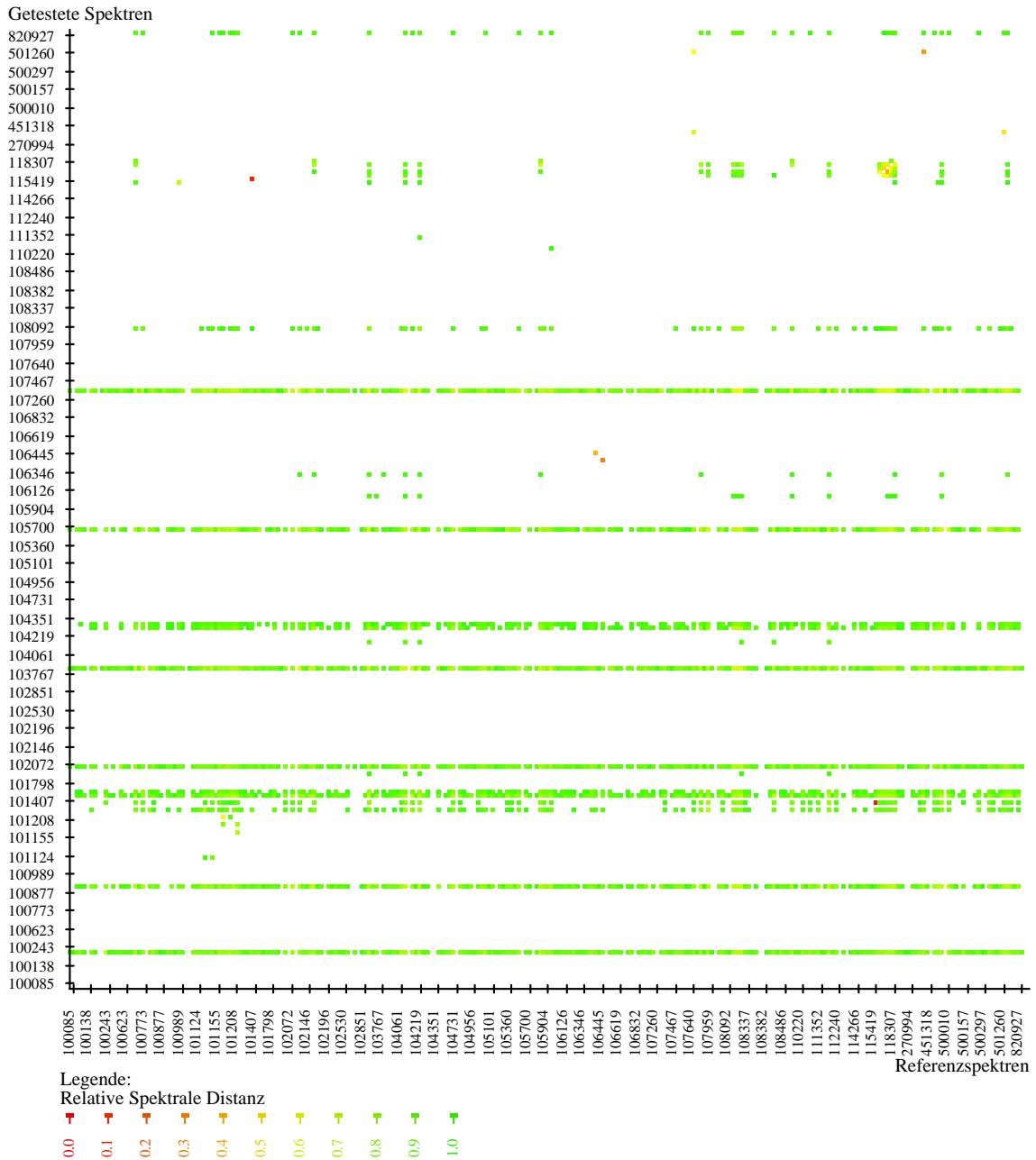
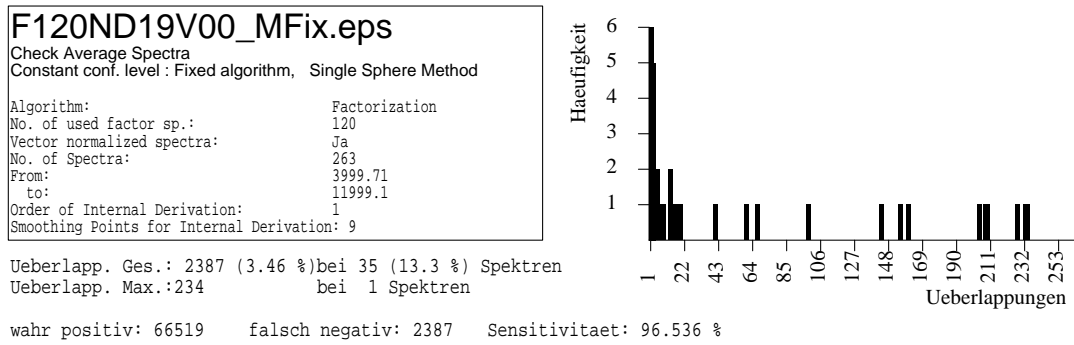


Abb. N17: Faktormethode (120 Faktoren, Vektornormierung und 1. Ableitung).

7.4.3 NIR, Variation des Spektralbereiches

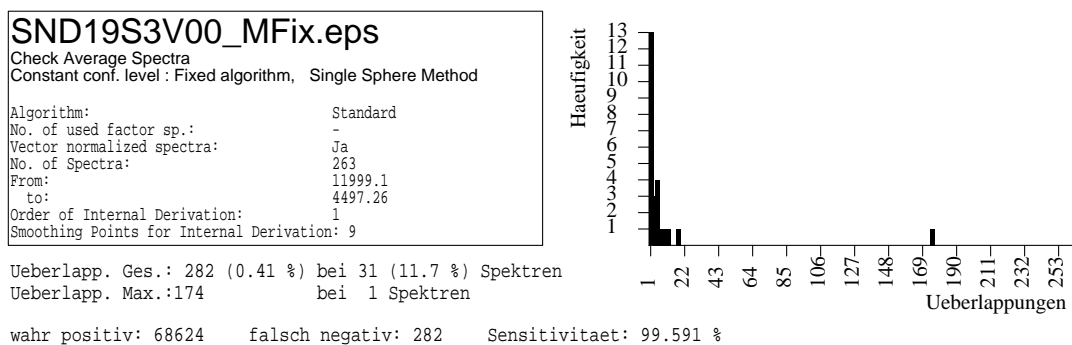


Abb. N18: Standardmethode (Vektornormierung und 1. Ableitung, 12000 cm^{-1} bis 4500 cm^{-1}).

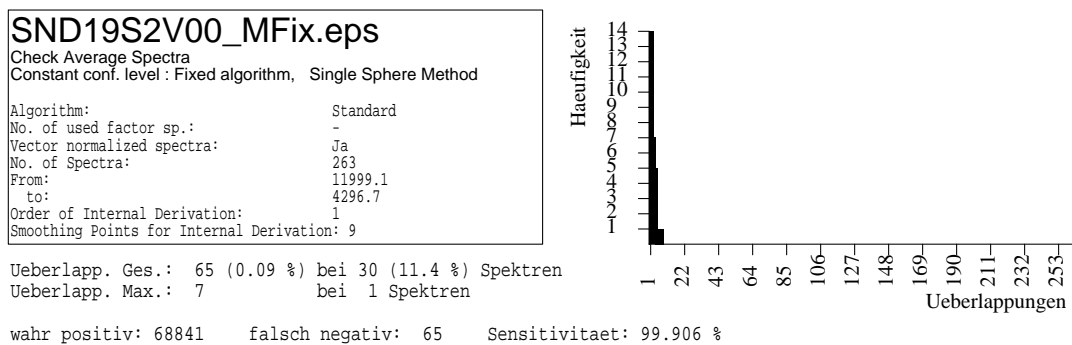


Abb. N19: Standardmethode (Vektornormierung und 1. Ableitung, 12000 cm^{-1} bis 4300 cm^{-1}).

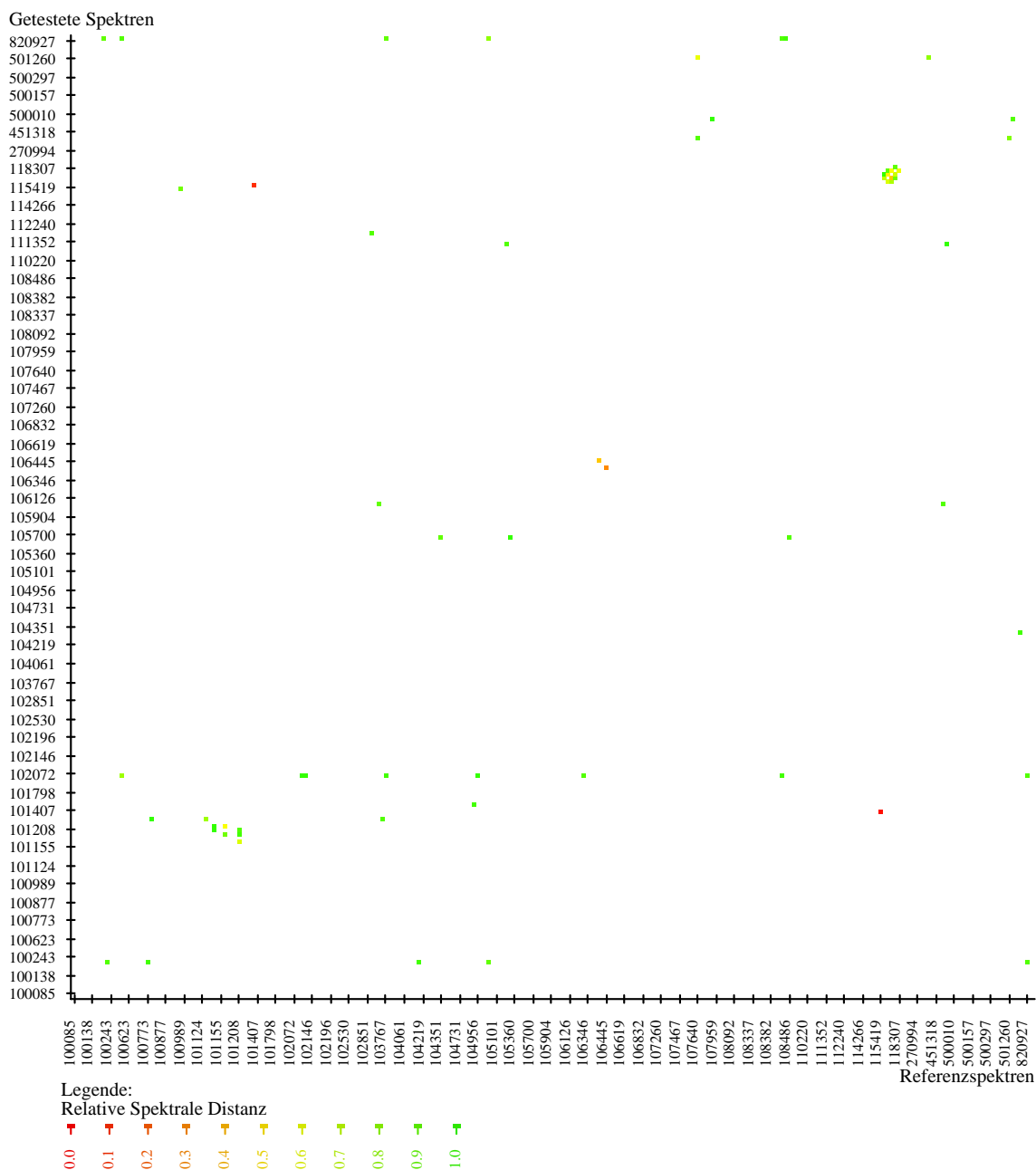
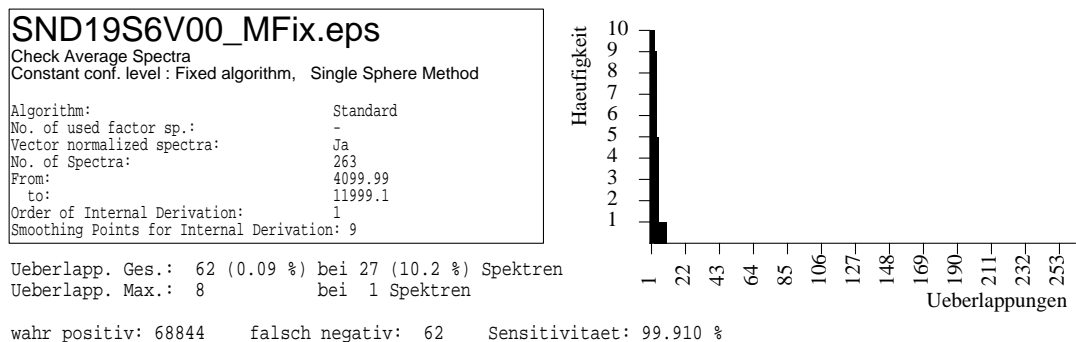


Abb. N20: Standardmethode (Vektornormierung und 1. Ableitung, 12000 cm^{-1} bis 4100 cm^{-1}).

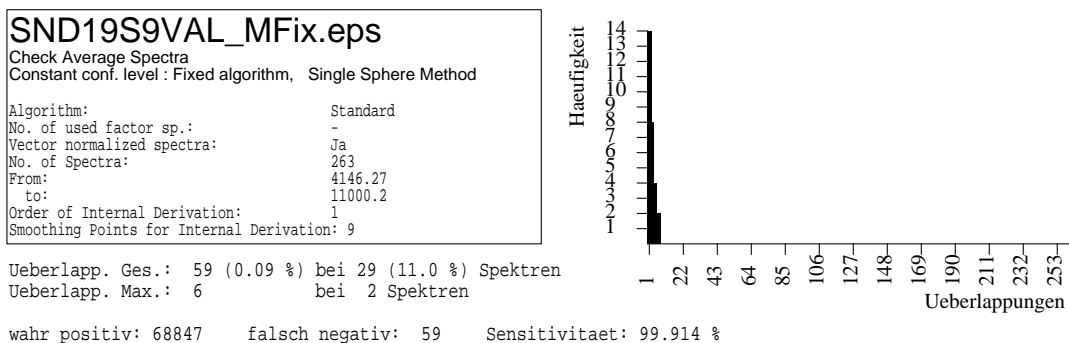


Abb. N21: Standardmethode (Vektornormierung und 1. Ableitung, 11000 cm^{-1} bis 4150 cm^{-1}).

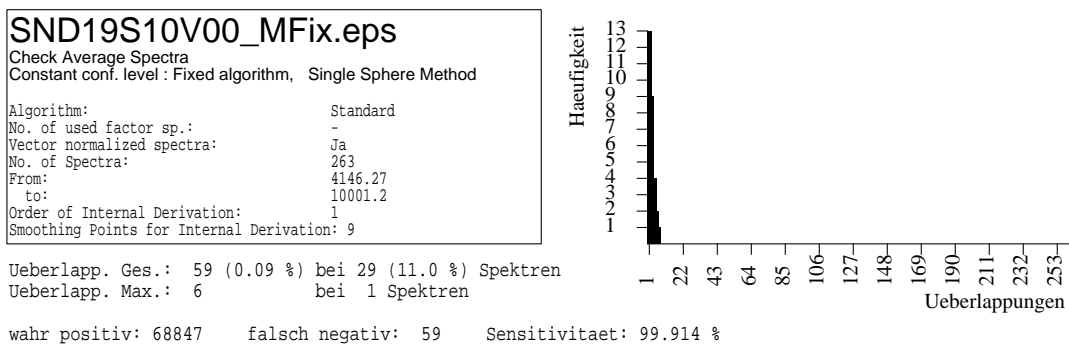


Abb. N22: Standardmethode (Vektornormierung und 1. Ableitung, 10000 cm^{-1} bis 4150 cm^{-1}).

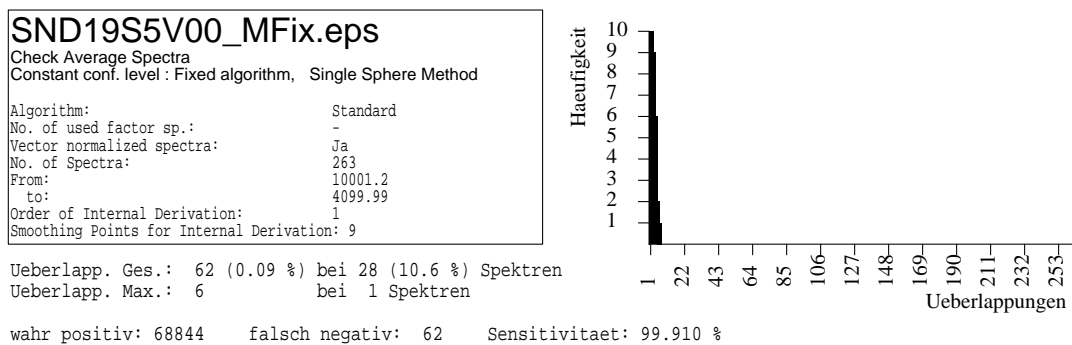


Abb. N23: Standardmethode (Vektornormierung und 1. Ableitung, 10000 cm^{-1} bis 4100 cm^{-1}).

7.4.4 NIR, Variation der Thresholdberechnung

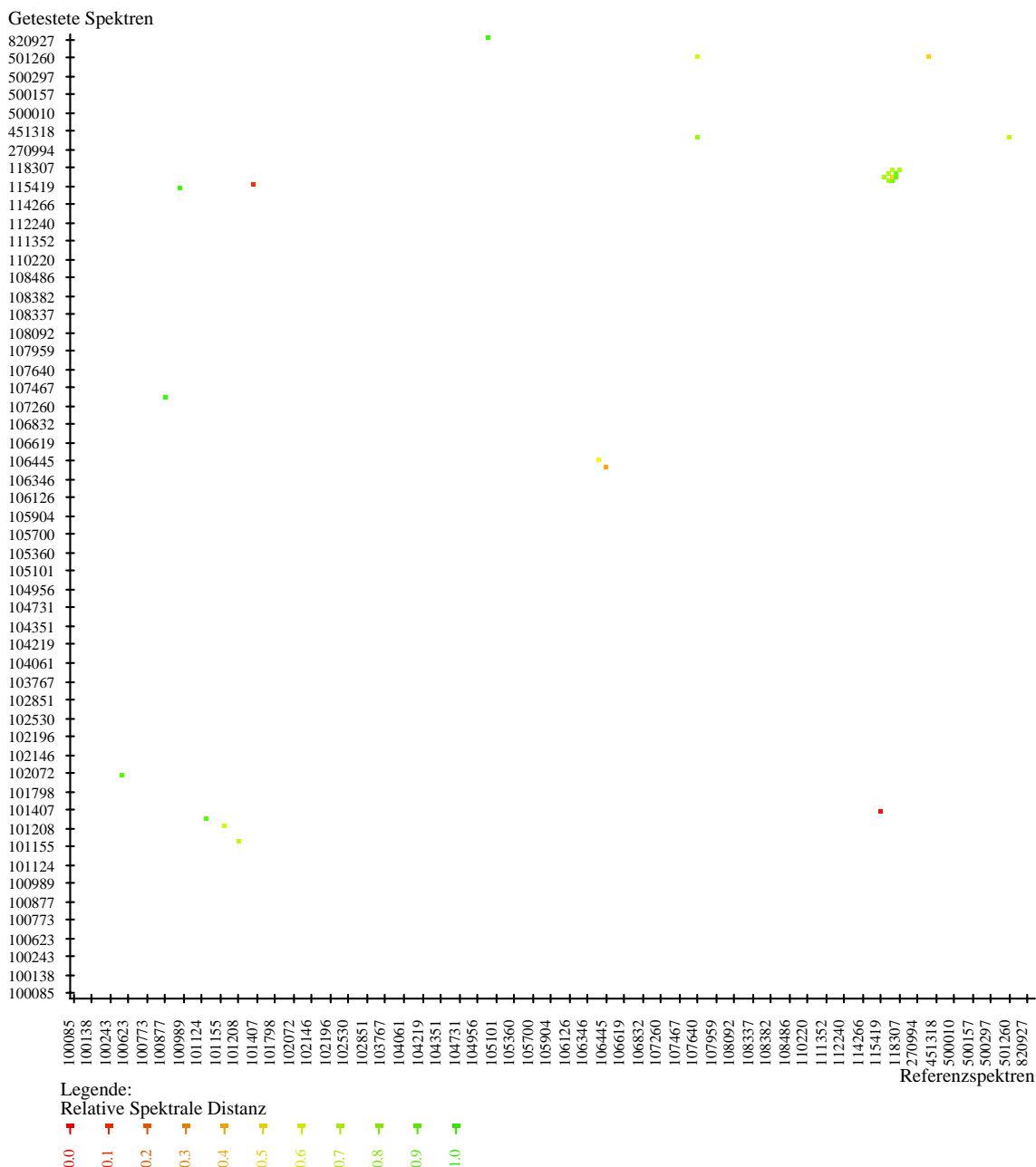
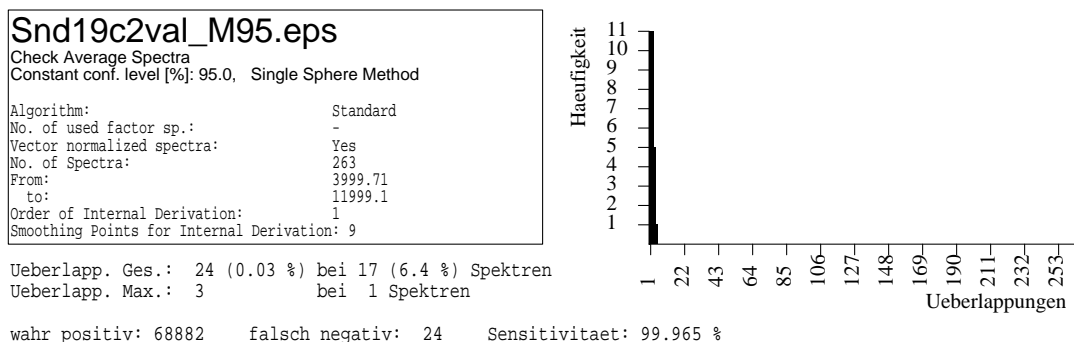
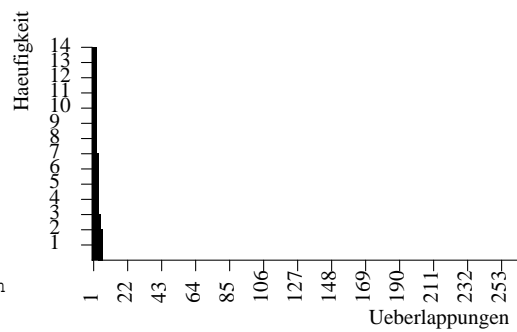


Abb. N24: Standardmethode (Vektornormierung und 1. Ableitung, konstantes Konfidenzniveau von 95 %).

SND19VAL_M99.eps
 Check Average Spectra
 Constant conf. level [%]: 99.0, Single Sphere Method

Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 To: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Überlapp. Ges.: 51 (0.07 %) bei 28 (10.6 %) Spektren
 Überlapp. Max.: 5 bei 2 Spektren

wahr positiv: 68855 falsch negativ: 51 Sensitivitaet: 99.926 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N25: Standardmethode (Vektornormierung und 1. Ableitung, konstantes Konfidenz-niveau von 99 %).

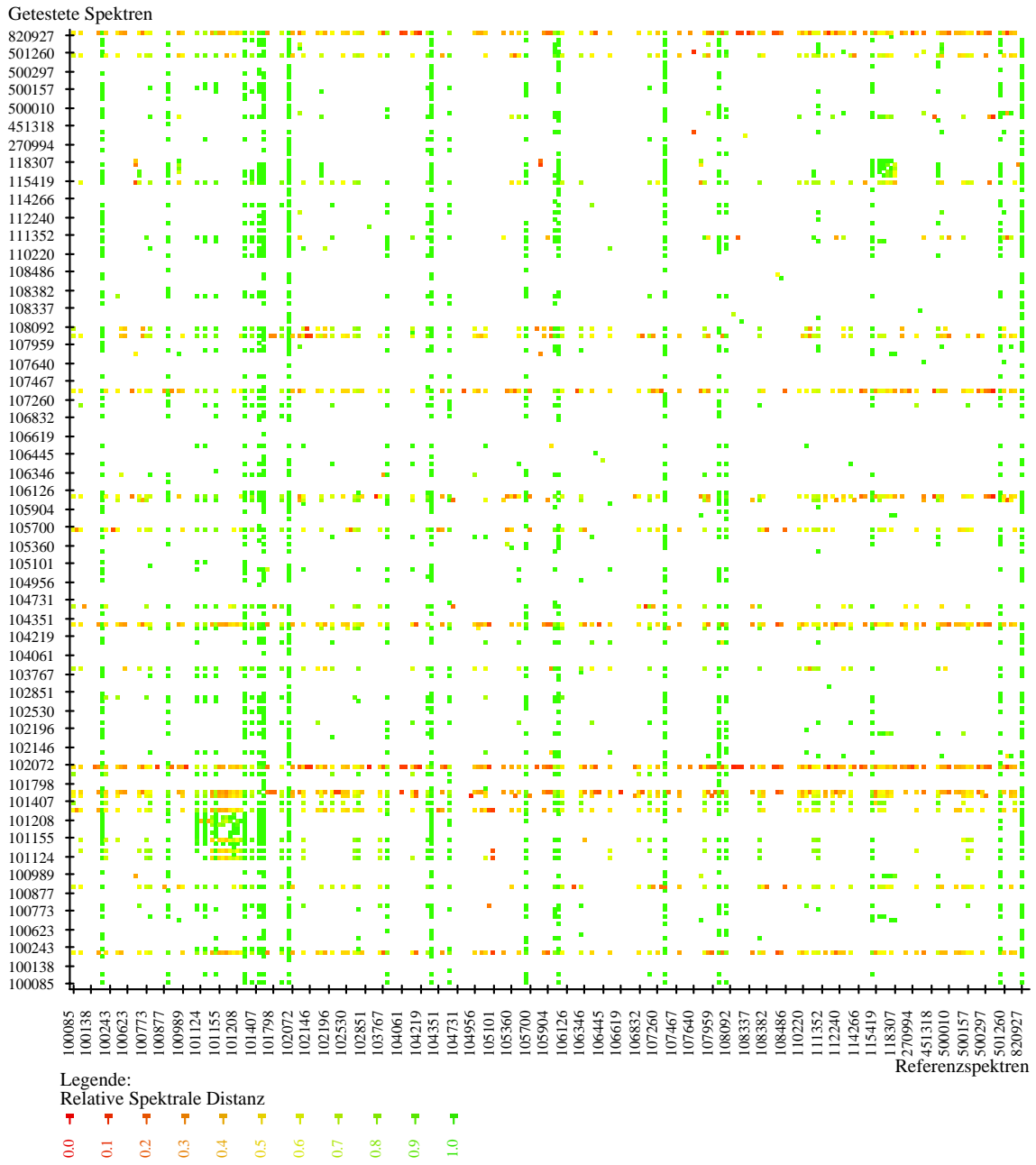
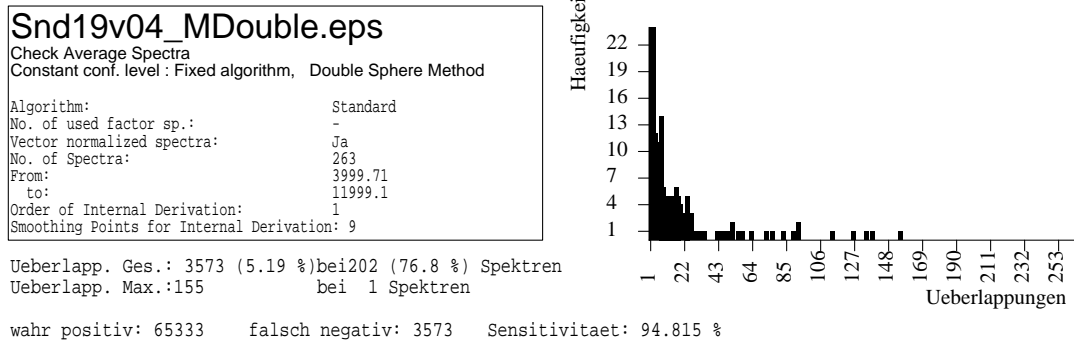
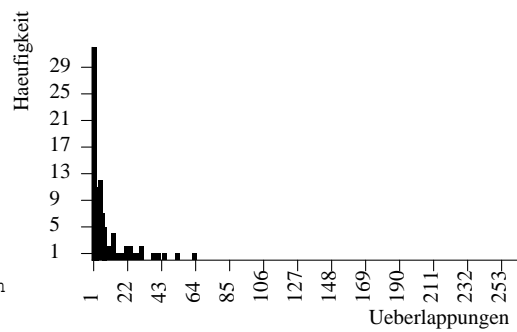


Abb. N26: Standardmethode (Vektornormierung und 1. Ableitung, Doppelkugelmodell, Fixed Algorithm).

SND19VAL_MDouble.eps
 Check Average Spectra
 Constant conf. level [%]: 95.0, Double Sphere Method

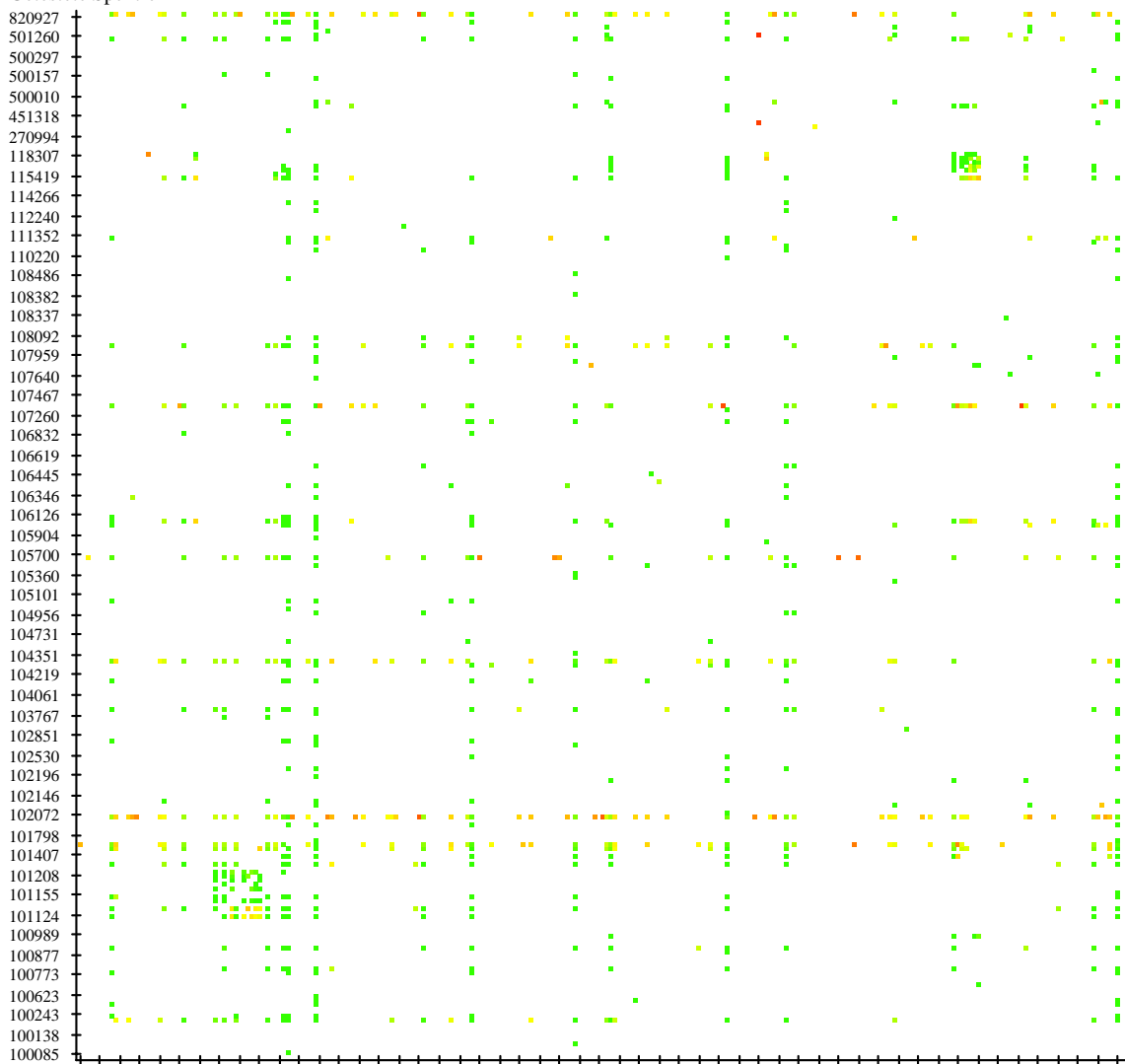
Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 898 (1.30 %) bei 115 (43.7 %) Spektren
 Ueberlapp. Max.: 63 bei 1 Spektren

wahr positiv: 68008 falsch negativ: 898 Sensitivitaet: 98.697 %

Getestete Spektren



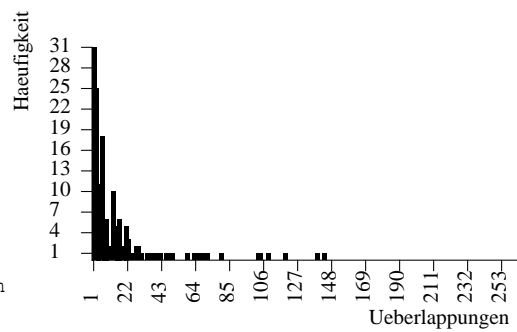
Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N27: Standardmethode (Vektornormierung und 1. Ableitung, Doppelkugelmodell, konstantes Konfidenzniveau von 95 %).

SND19bVAL_MDouble.eps
 Check Average Spectra
 Constant conf. level [%]: 99.0, Double Sphere Method

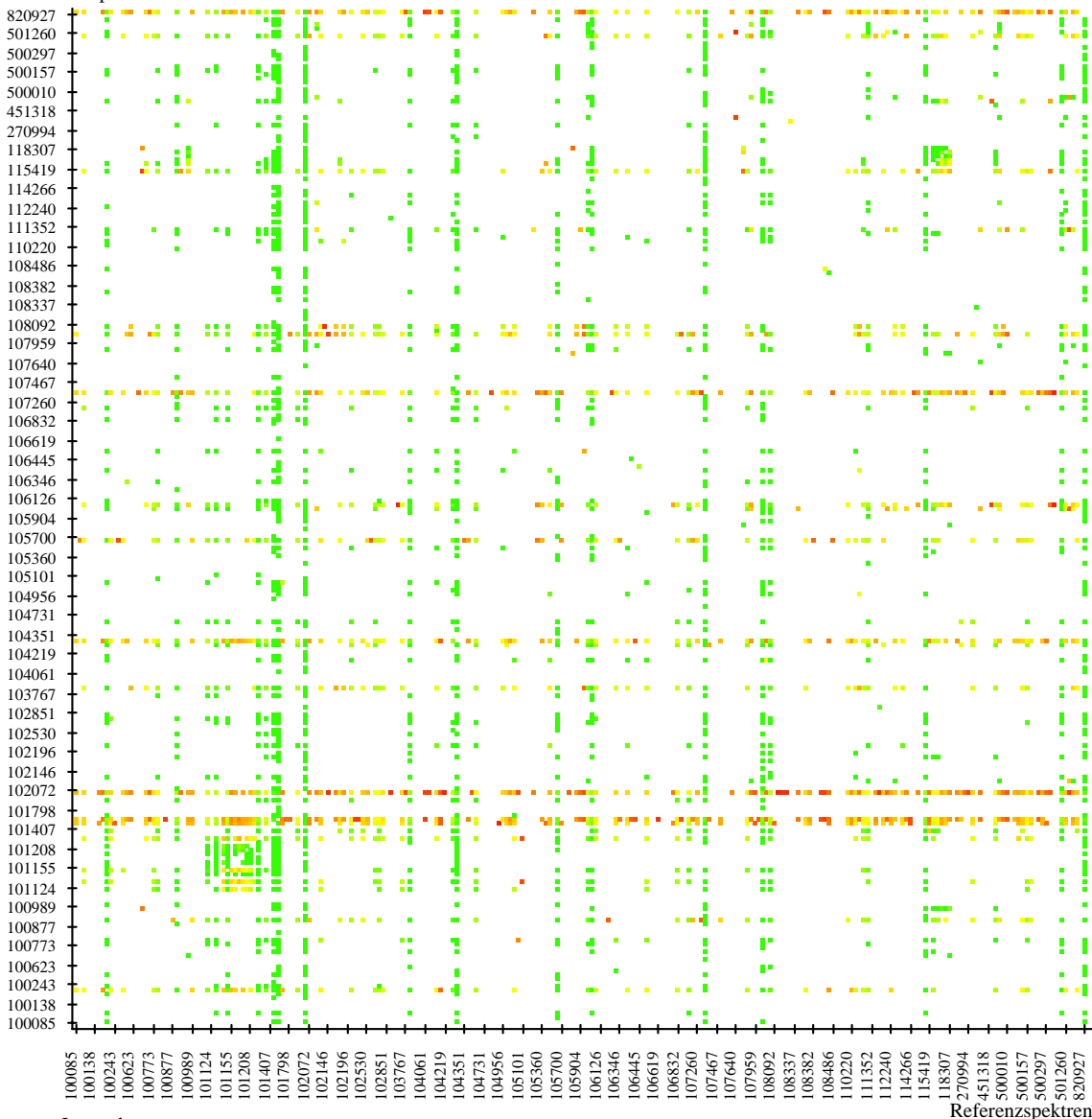
Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 To: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 2988 (4.34 %) bei 196 (74.5 %) Spektren
 Ueberlapp. Max.: 143 bei 1 Spektren

wahr positiv: 65918 falsch negativ: 2988 Sensitivitaet: 95.664 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. N28: Standardmethode (Vektornormierung und 1. Ableitung, Doppelkugelmodell, konstantes Konfidenzniveau von 99 %).

7.4.5 Raman, Standardmethode

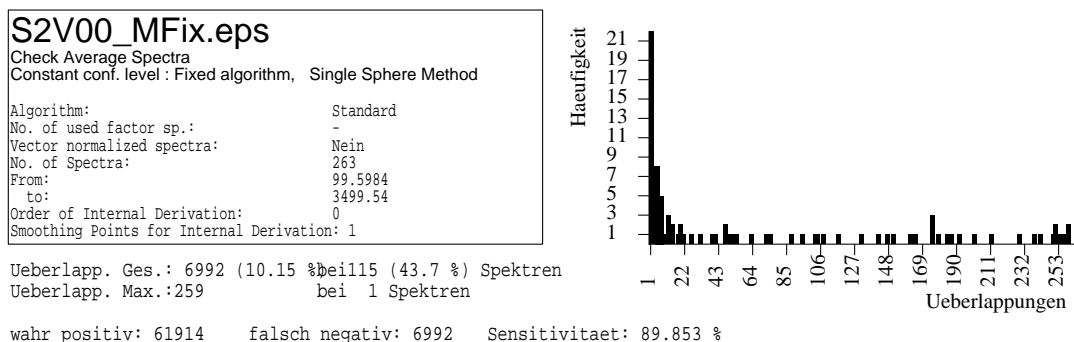
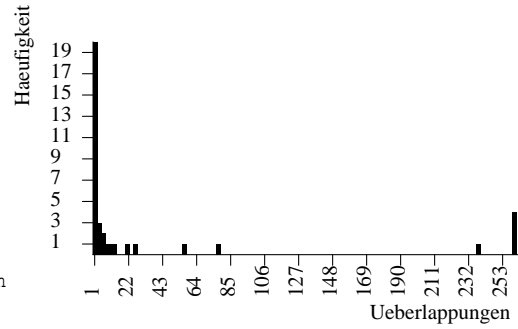


Abb. R1: Standardmethode.

```

snavl_MFix.eps
Check Average Spectra
Constant conf. level : Fixed algorithm, Single Sphere Method

Algorithm:                               Standard
No. of used factor sp.:                  -
Vector normalized spectra:               Ja
No. of Spectra:                          263
From:                                    99.5984
to:                                       3499.54
Order of Internal Derivation:             0
Smoothing Points for Internal Derivation: 1
    
```



Ueberlapp. Ges.: 1555 (2.26 %) bei 44 (16.7 %) Spektren
 Ueberlapp. Max.: 260 bei 4 Spektren

wahr positiv: 67351 falsch negativ: 1555 Sensitivitaet: 97.743 %

Getestete Spektren



Abb. R2: Standardmethode (Vektornormierung).

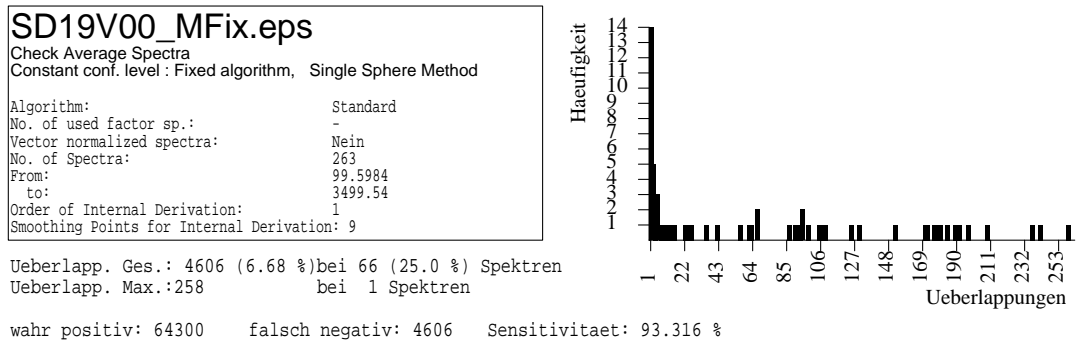


Abb. R3: Standardmethode (1. Ableitung).

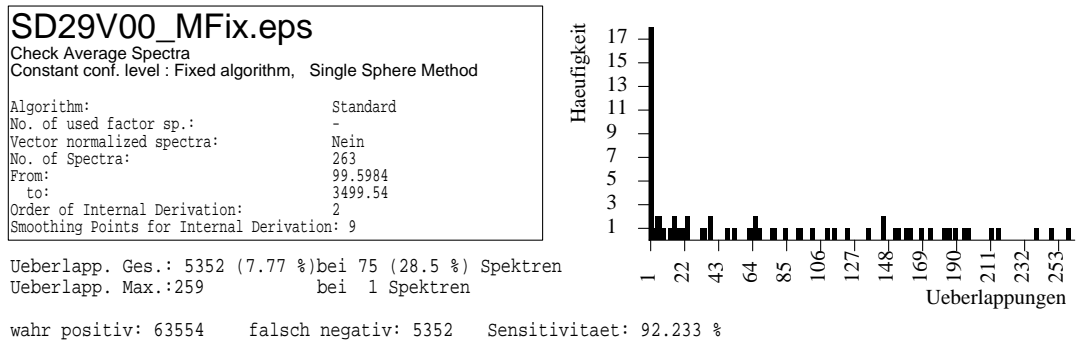


Abb. R4: Standardmethode (2. Ableitung).

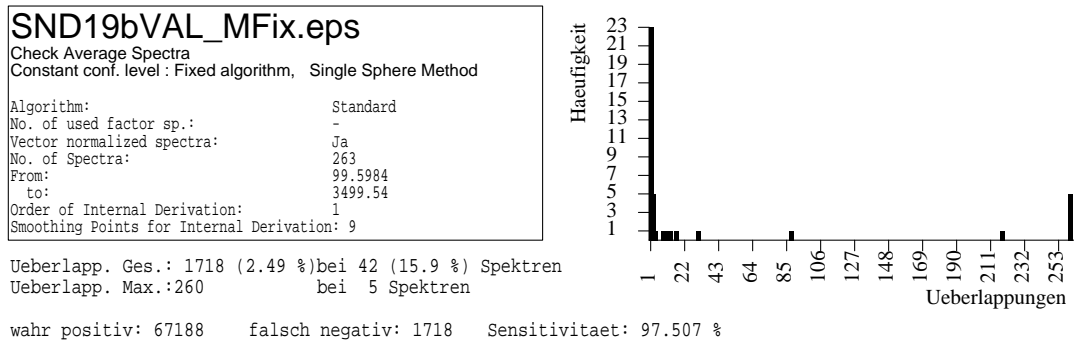


Abb. R5: Standardmethode (Vektornormierung und 1. Ableitung).

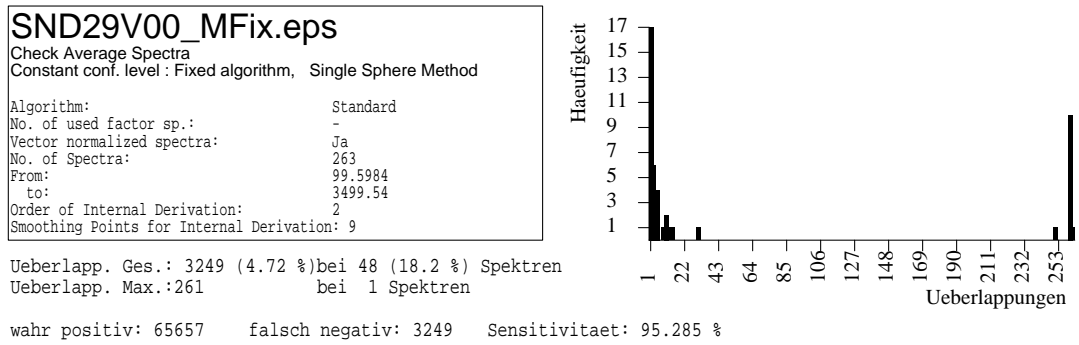
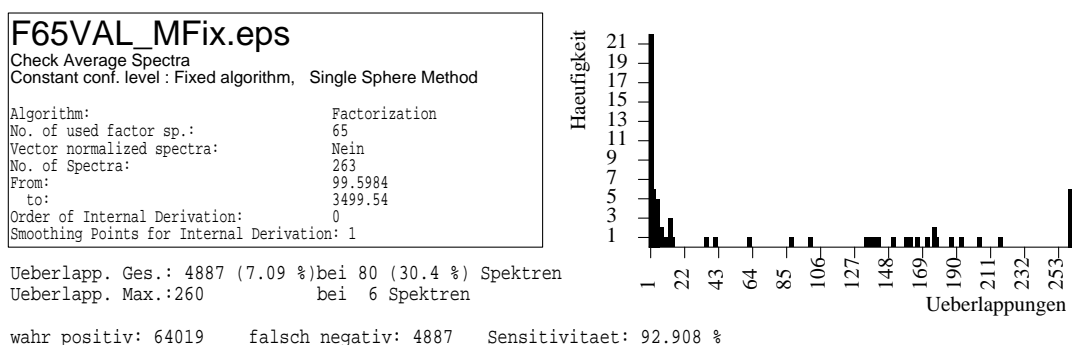


Abb. R6: Standardmethode (Vektornormierung und 2. Ableitung).

7.4.6 Raman, Faktormethode



Getestete Spektren

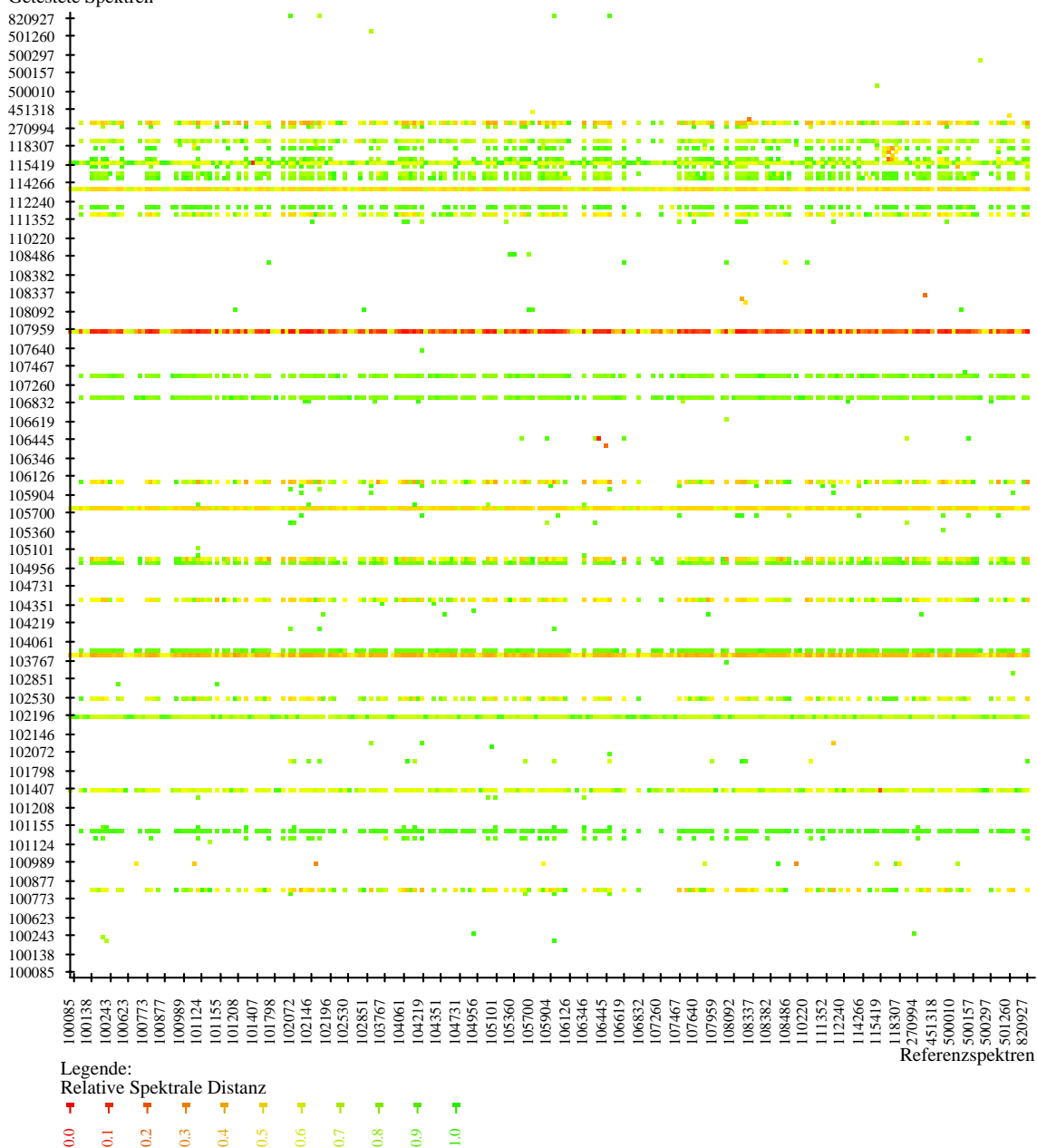
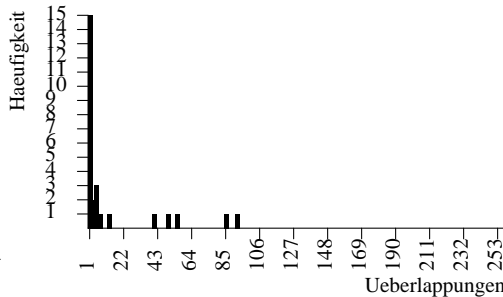


Abb. R7: Faktormethode (65 Faktoren).

F65NVAL_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 99.5984
 to: 3499.54
 Order of Internal Derivation: 0
 Smoothing Points for Internal Derivation: 1



Ueberlapp. Ges.: 1430 (2.08 %) bei 36 (13.6 %) Spektren
 Ueberlapp. Max.: 260 bei 4 Spektren

wahr positiv: 67476 falsch negativ: 1430 Sensitivitaet: 97.925 %

Getestete Spektren

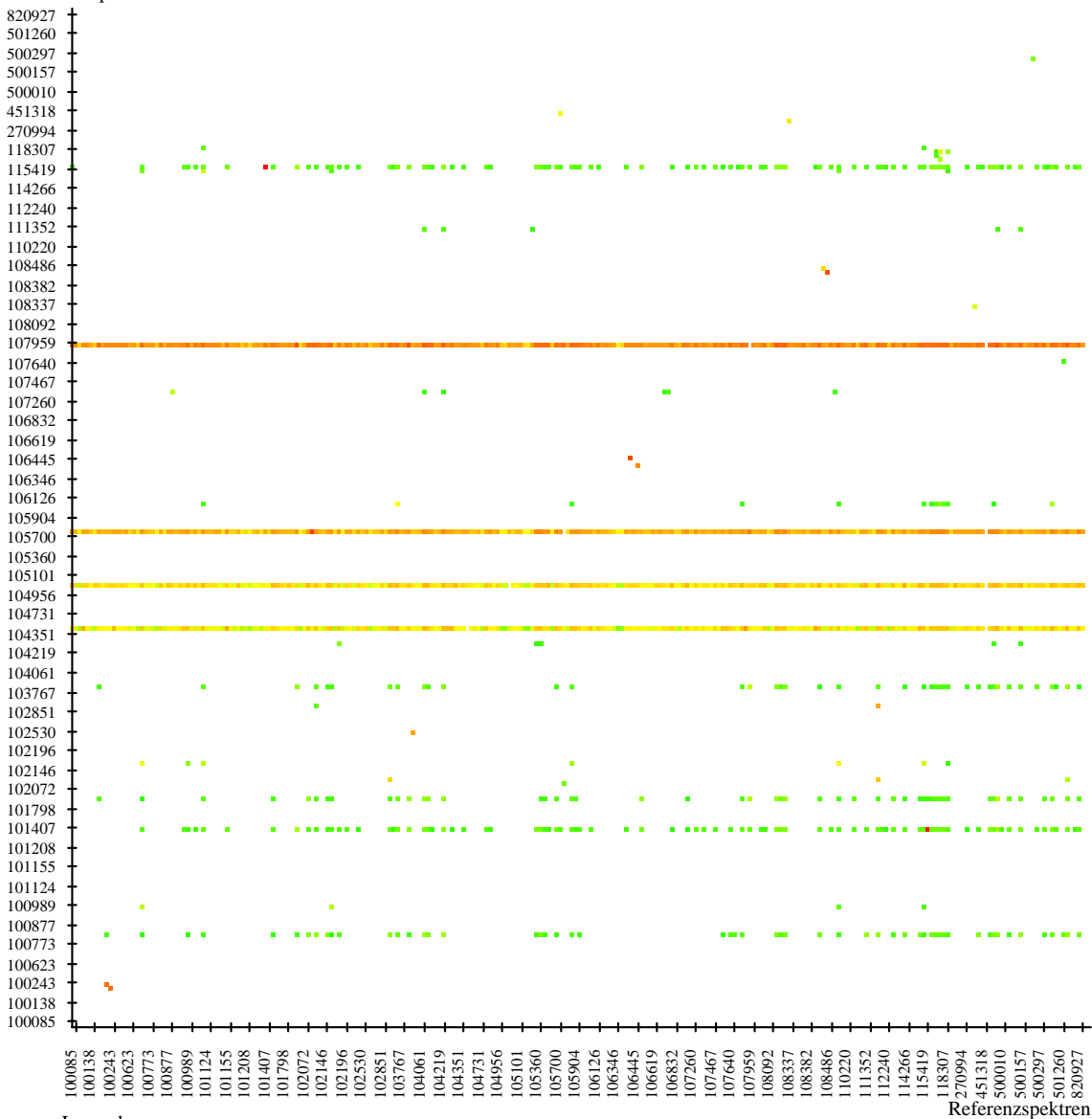


Abb. R8: Faktormethode (65 Faktoren, Vektornormierung).

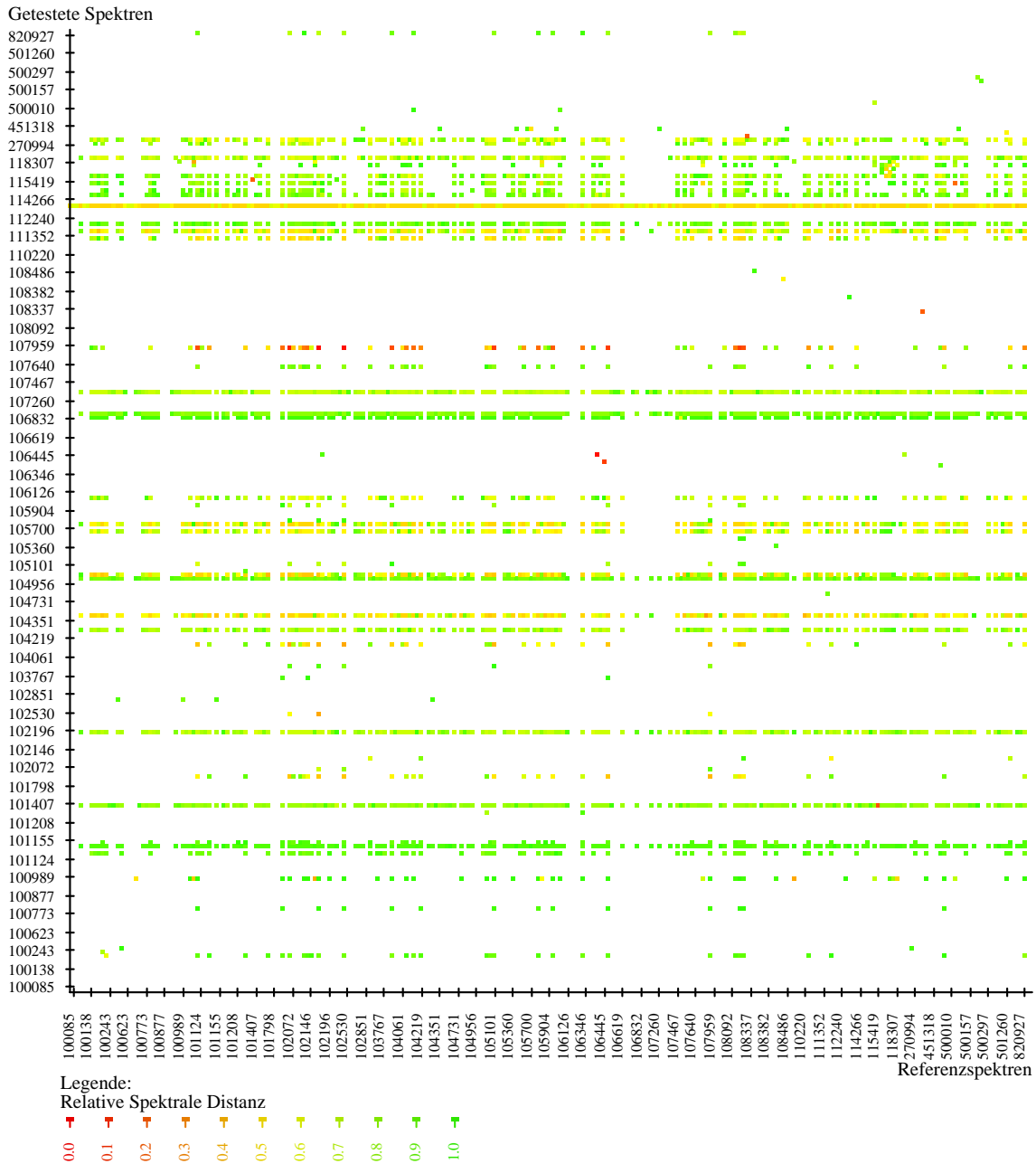
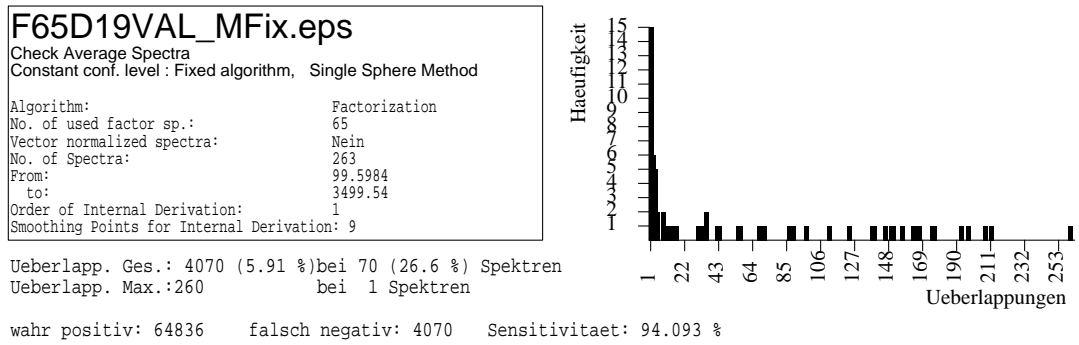
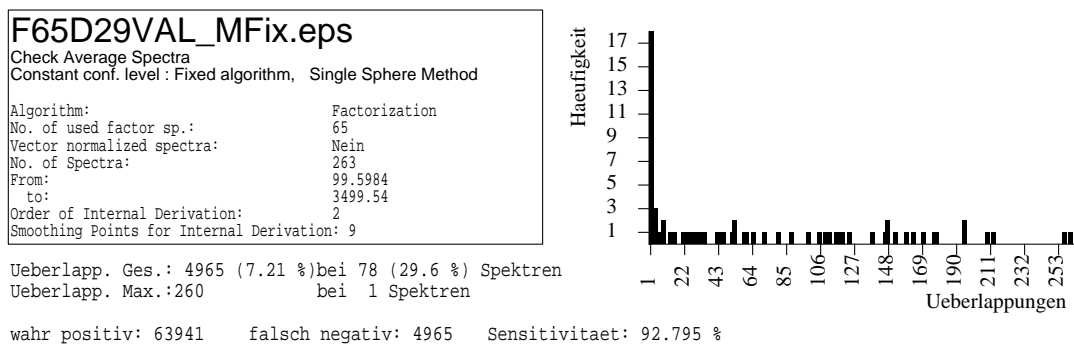


Abb. R9: Faktormethode (65 Faktoren, 1. Ableitung).



Getestete Spektren

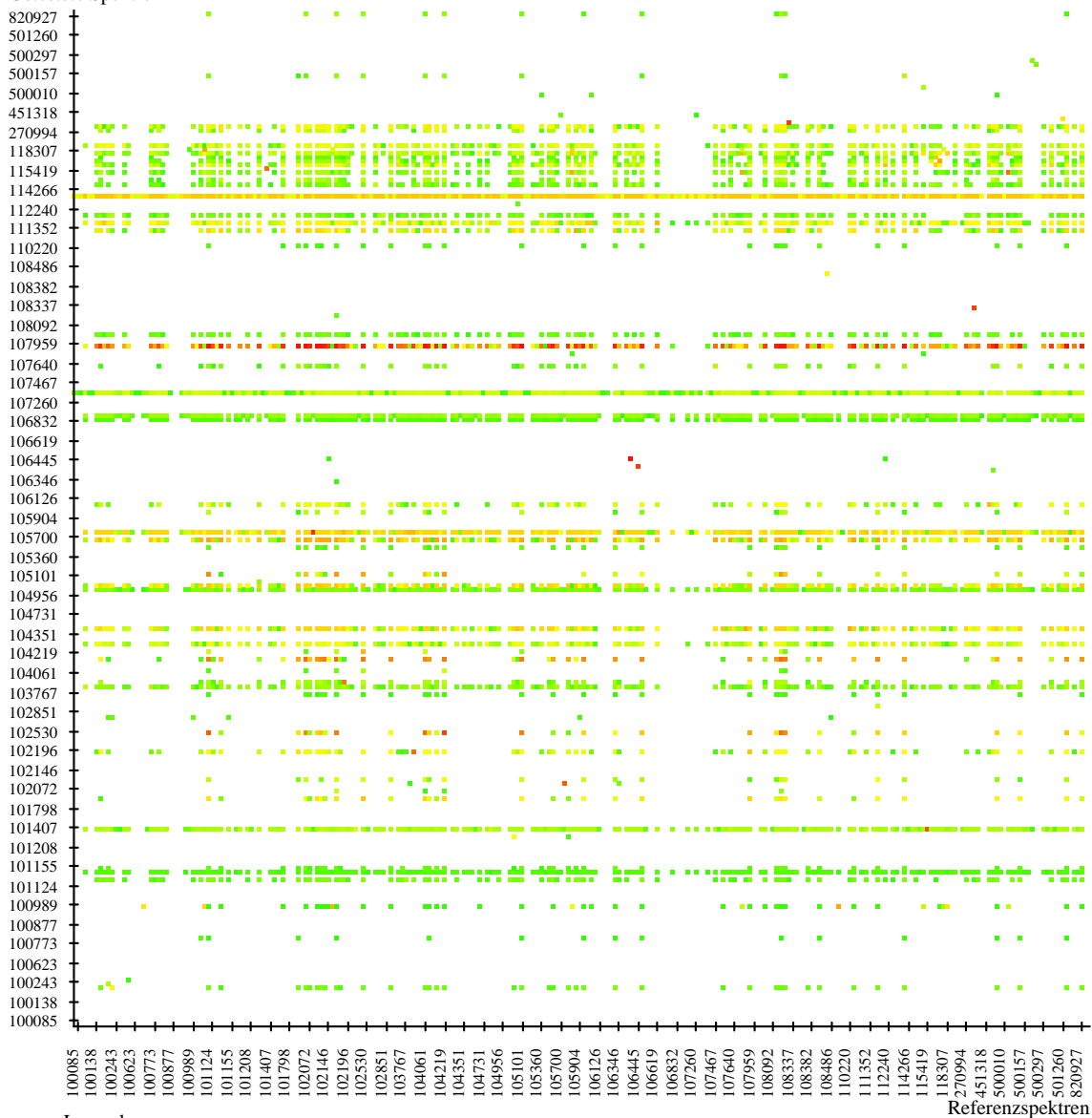
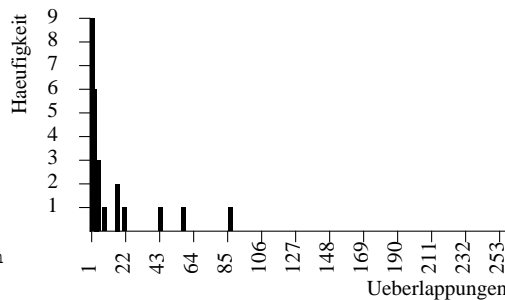


Abb. R10: Faktormethode (65 Faktoren, 2. Ableitung).

F65ND19V00_MFix.eps
 Check Average Spectra
 Constant conf. level : Fixed algorithm, Single Sphere Method

Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 to: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 292 (0.42 %) bei 28 (10.6 %) Spektren
 Ueberlapp. Max.: 86 bei 1 Spektren

wahr positiv: 68614 falsch negativ: 292 Sensitivitaet: 99.576 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

- 0.0
- 0.1
- 0.2
- 0.3
- 0.4
- 0.5
- 0.6
- 0.7
- 0.8
- 0.9
- 1.0

Abb. R11: Faktormethode (65 Faktoren, Vektornormierung, 1. Ableitung).

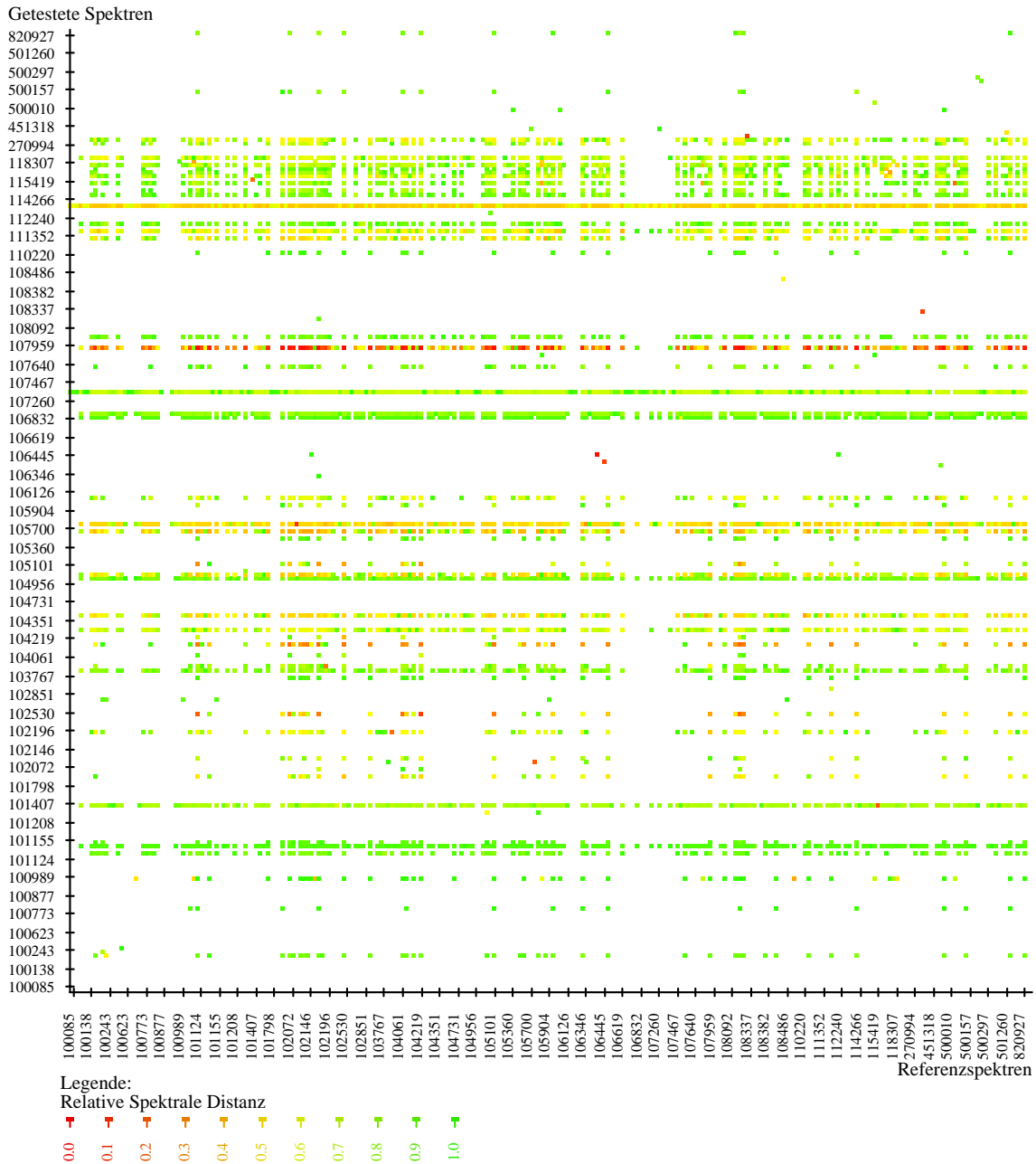
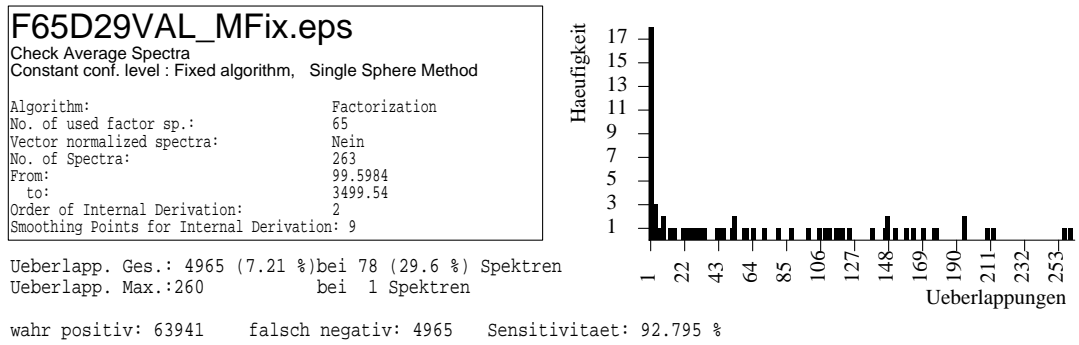


Abb. R12: Faktormethode (65 Faktoren, Vektornormierung, 2. Ableitung).

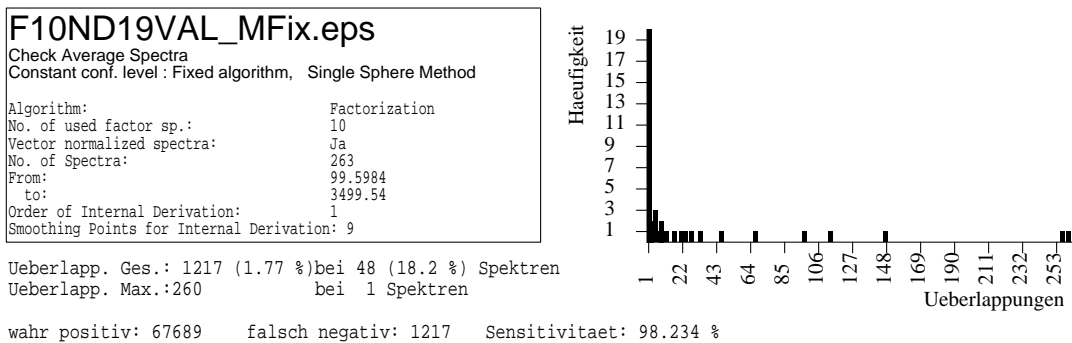
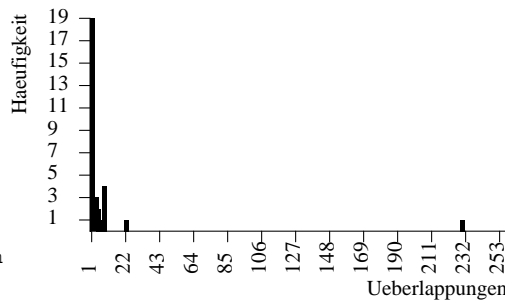


Abb. R13: Faktormethode (10 Faktoren, Vektornormierung, 1. Ableitung).

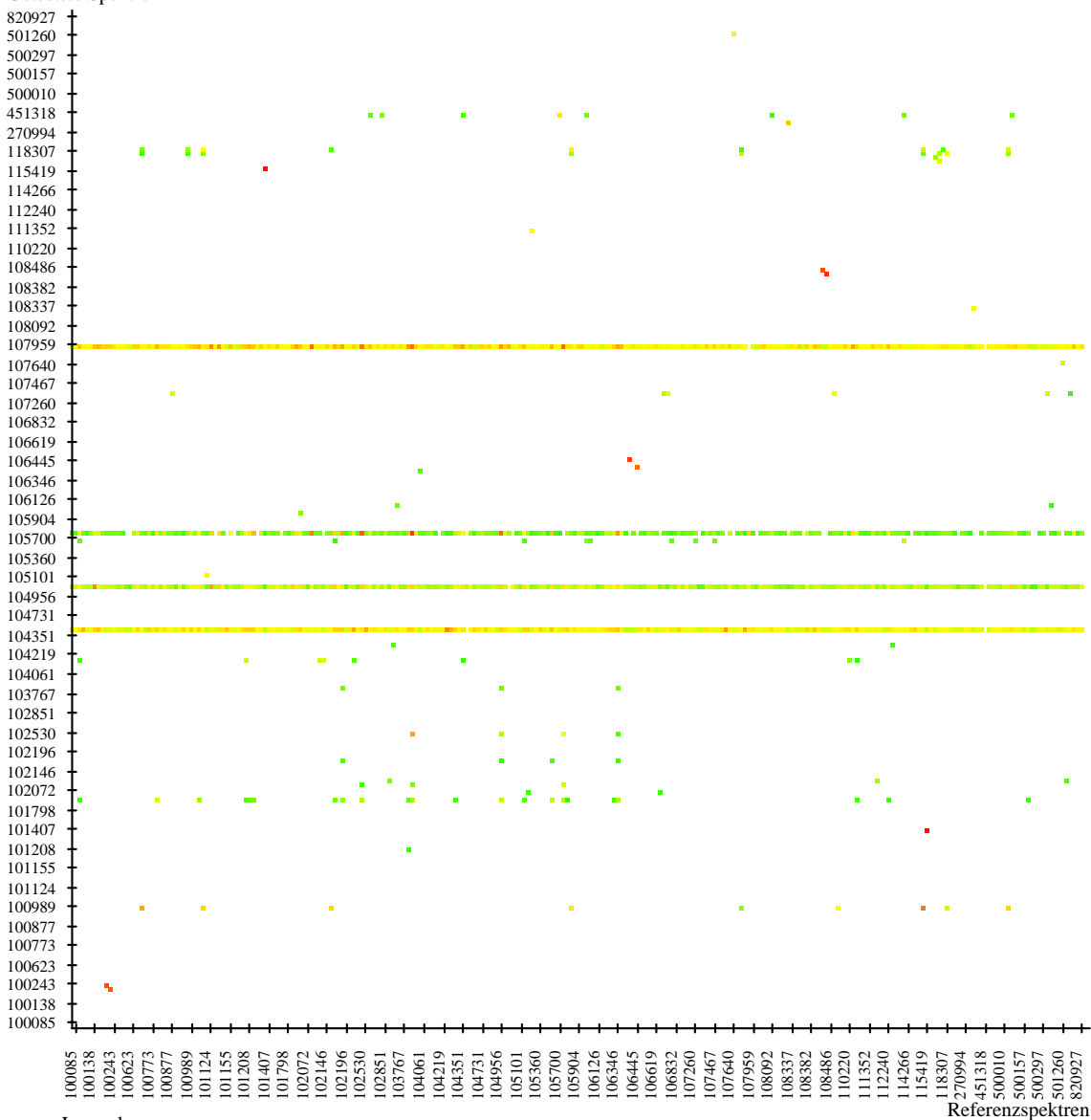
F25ND19VAL_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method
 Algorithm: Factorization
 No. of used factor sp.: 25
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 99.5984
 to: 3499.54
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 1132 (1.64 %) bei 39 (14.8 %) Spektren
 Ueberlapp. Max.: 260 bei 3 Spektren

wahr positiv: 67774 falsch negativ: 1132 Sensitivitaet: 98.357 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz
 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. R14: Faktormethode (25 Faktoren, Vektornormierung, 1. Ableitung).

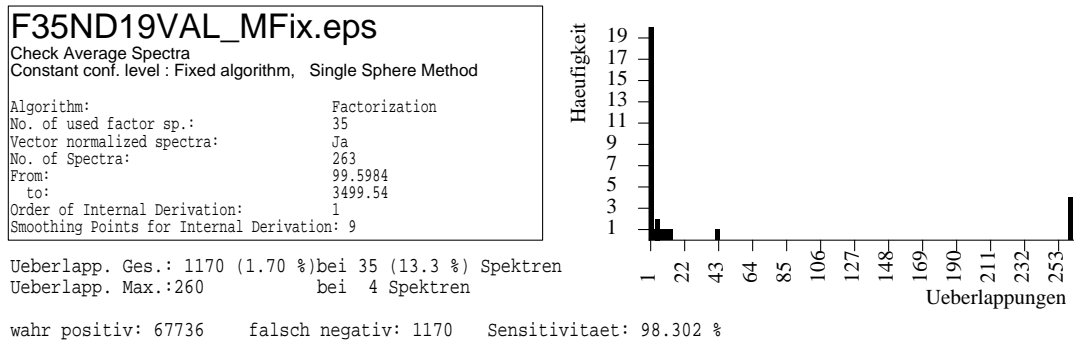
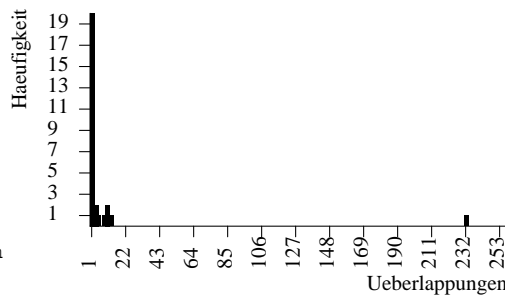


Abb. R15: Faktormethode (35 Faktoren, Vektornormierung, 1. Ableitung).

F45ND19VAL_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method
 Algorithm: Factorization
 No. of used factor sp.: 45
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 99.5984
 to: 3499.54
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 1348 (1.96 %) bei 34 (12.9 %) Spektren
 Ueberlapp. Max.: 260 bei 4 Spektren

wahr positiv: 67558 falsch negativ: 1348 Sensitivitaet: 98.044 %

Getestete Spektren



Abb. R16: Faktormethode (45 Faktoren, Vektornormierung, 1. Ableitung).

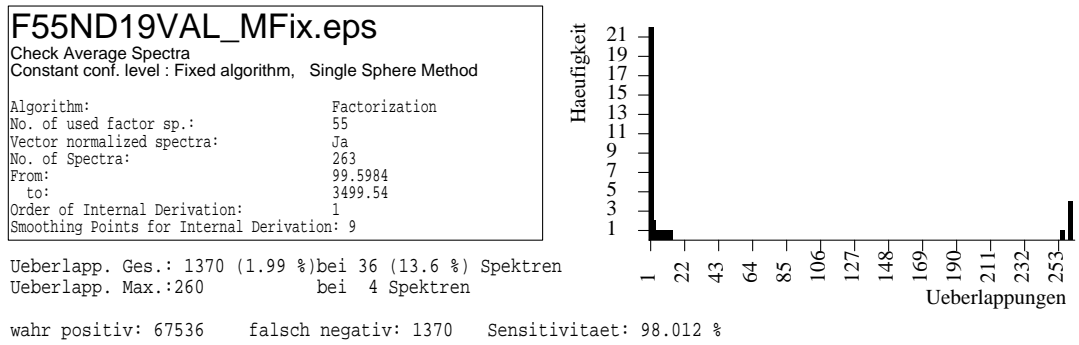
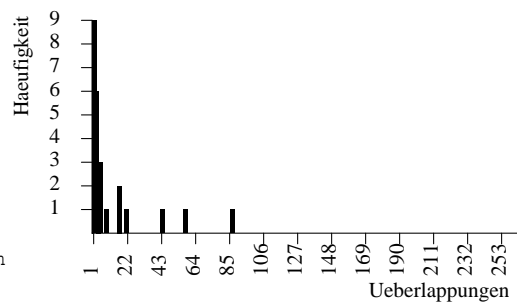


Abb. R17: Faktormethode (55 Faktoren, Vektornormierung, 1. Ableitung).

F65ND19V00_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method

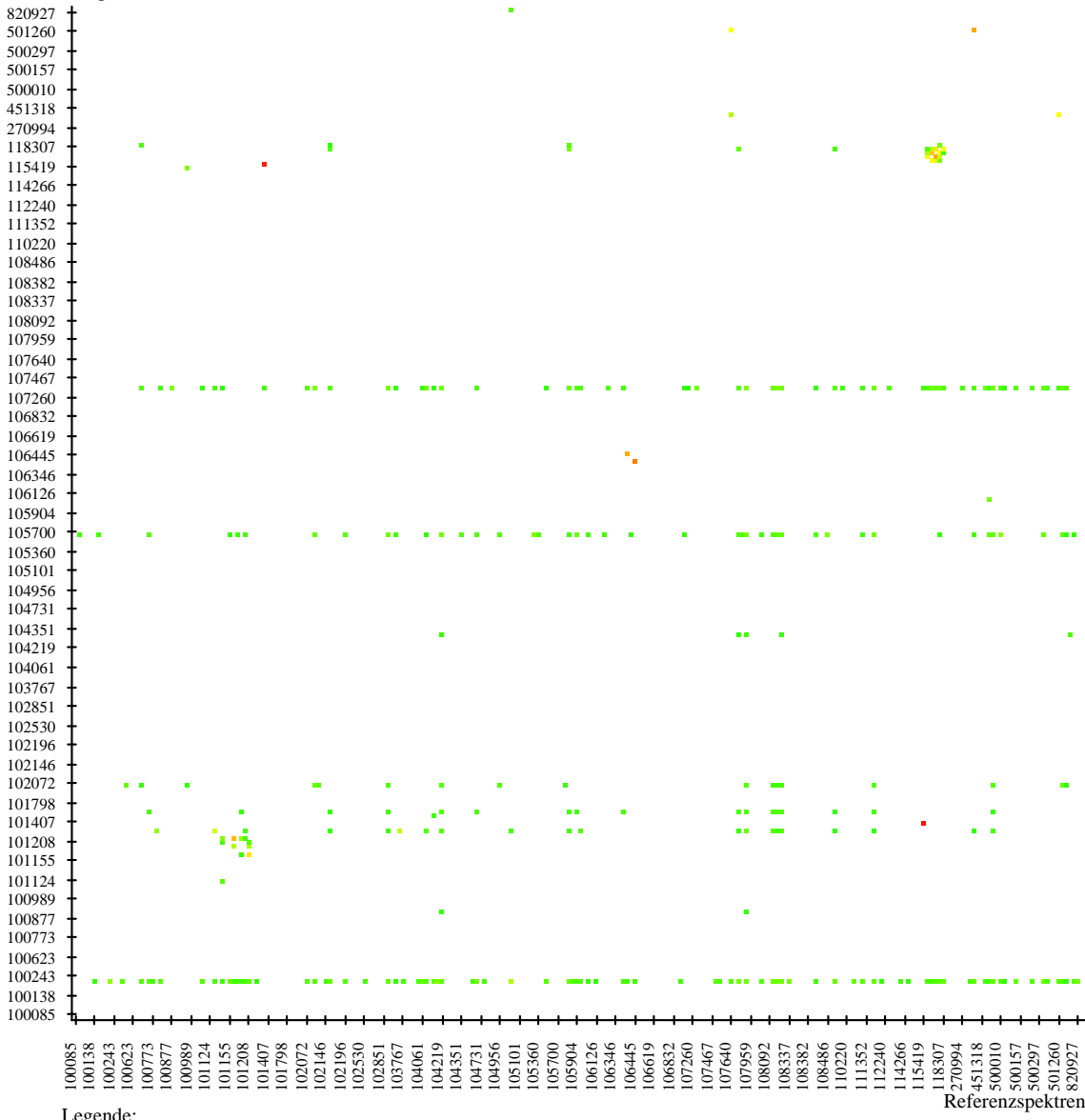
Algorithm: Factorization
 No. of used factor sp.: 65
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 3999.71
 To: 11999.1
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 292 (0.42 %) bei 28 (10.6 %) Spektren
 Ueberlapp. Max.: 86 bei 1 Spektren

wahr positiv: 68614 falsch negativ: 292 Sensitivitaet: 99.576 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 (red square)
 0.1 (orange square)
 0.2 (light orange square)
 0.3 (yellow square)
 0.4 (light green square)
 0.5 (green square)
 0.6 (darker green square)
 0.7 (medium green square)
 0.8 (darker green square)
 0.9 (medium green square)
 1.0 (darker green square)

Abb. R18: Faktormethode (65 Faktoren, Vektornormierung, 1. Ableitung).

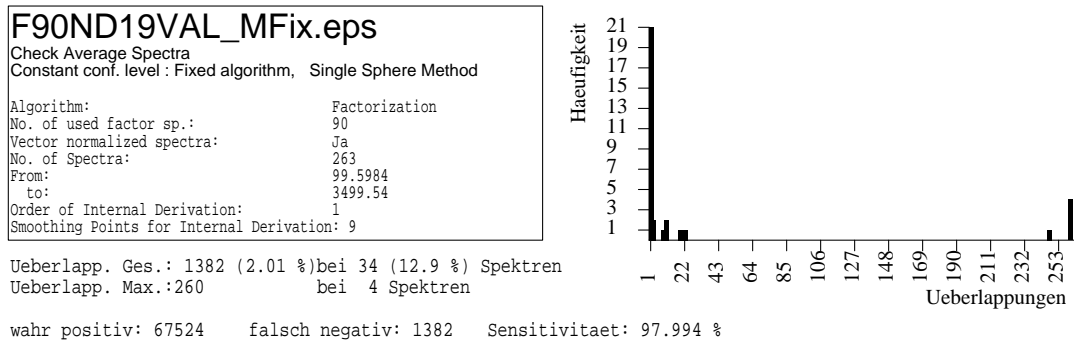
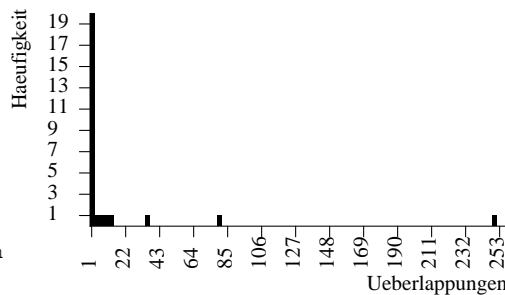


Abb. R19: Faktormethode (90 Faktoren, Vektornormierung, 1. Ableitung).

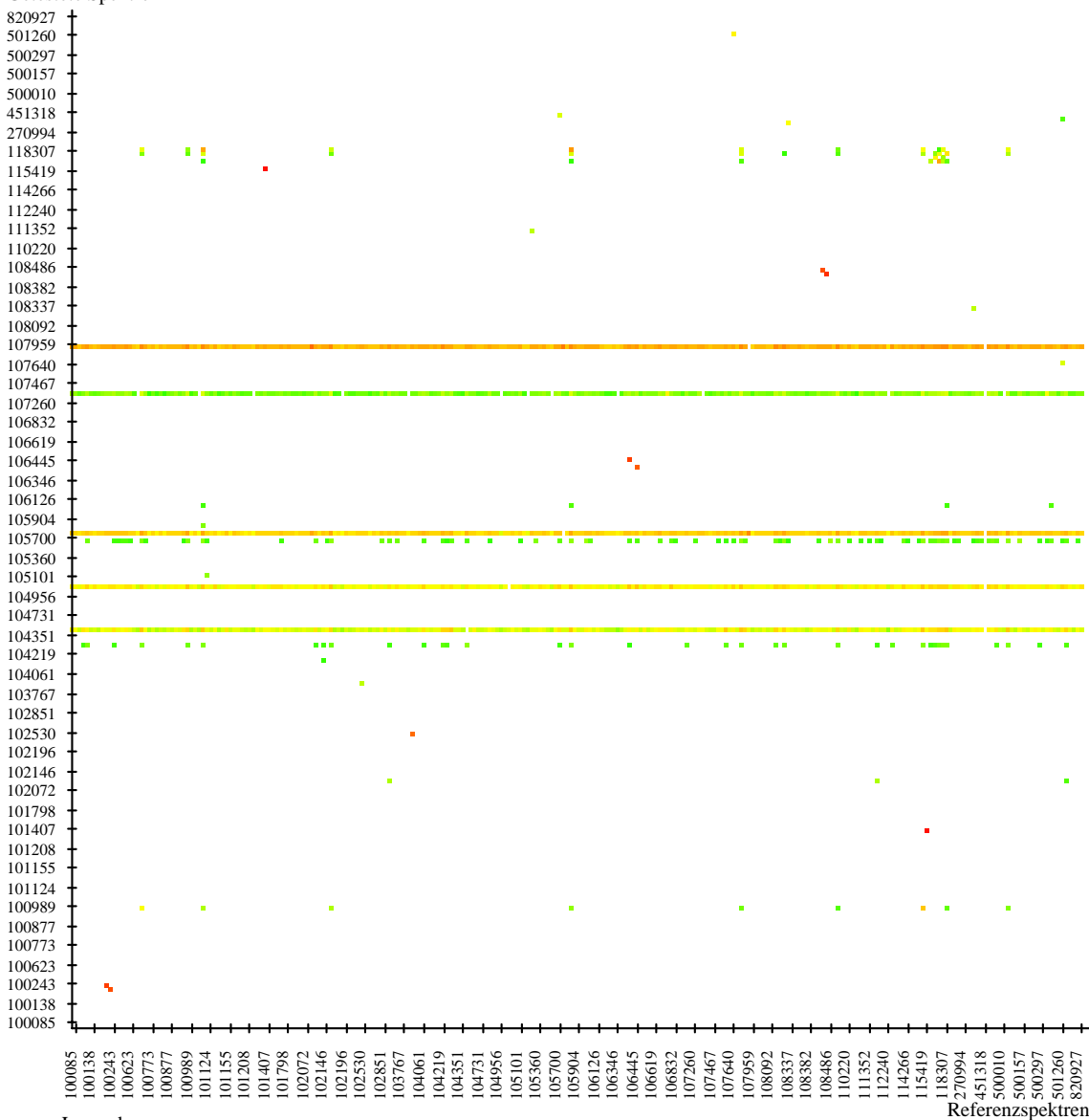
F120ND19VAL_MFix.eps
 Check Average Spectra
 Constant conf. level: Fixed algorithm, Single Sphere Method
 Algorithm: Factorization
 No. of used factor sp.: 120
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 99.5984
 to: 3499.54
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 1472 (2.14 %) bei 35 (13.3 %) Spektren
 Ueberlapp. Max.: 260 bei 4 Spektren

wahr positiv: 67434 falsch negativ: 1472 Sensitivitaet: 97.864 %

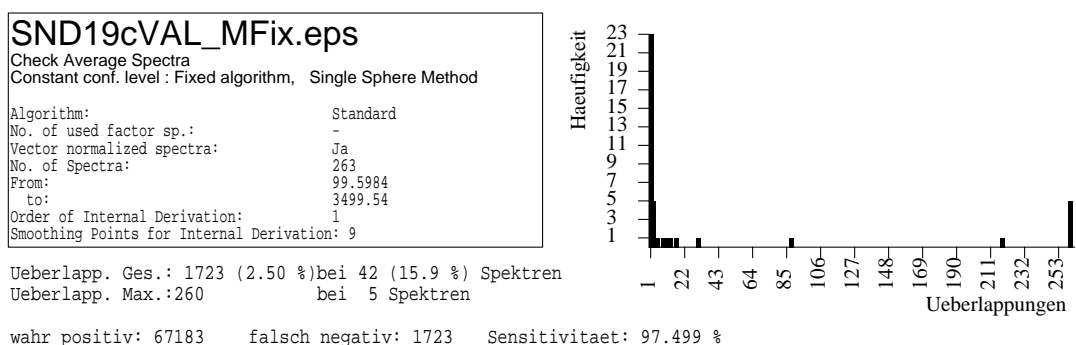
Getestete Spektren



Legende:
 Relative Spektrale Distanz
 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. R20: Faktormethode (120 Faktoren, Vektornormierung, 1. Ableitung).

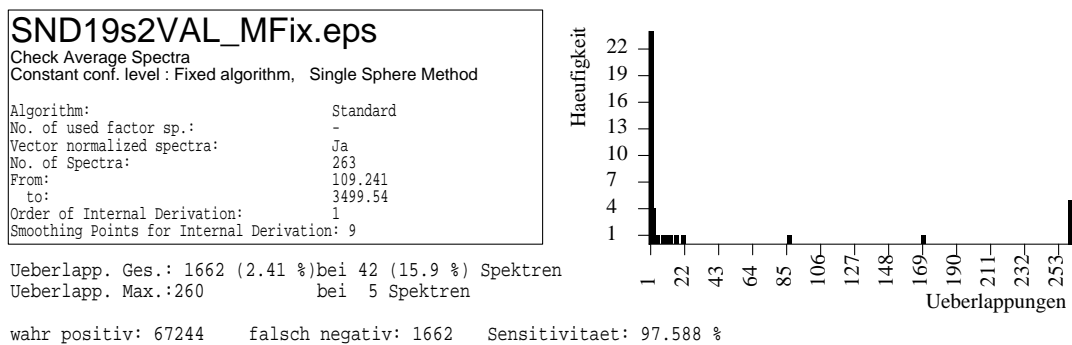
7.4.7 Raman, Variation des Spektralbereiches



Getestete Spektren



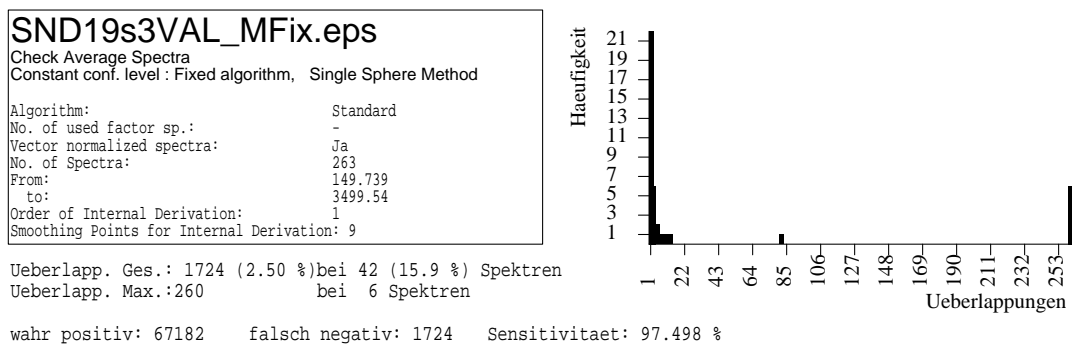
Abb. R21: Standardmethode (Vektornormierung, 1. Ableitung, 3500 cm^{-1} bis 100 cm^{-1}).



Getestete Spektren



Abb. R22: Standardmethode (Vektornormierung, 1. Ableitung, 3500 cm^{-1} bis 110 cm^{-1}).



Getestete Spektren

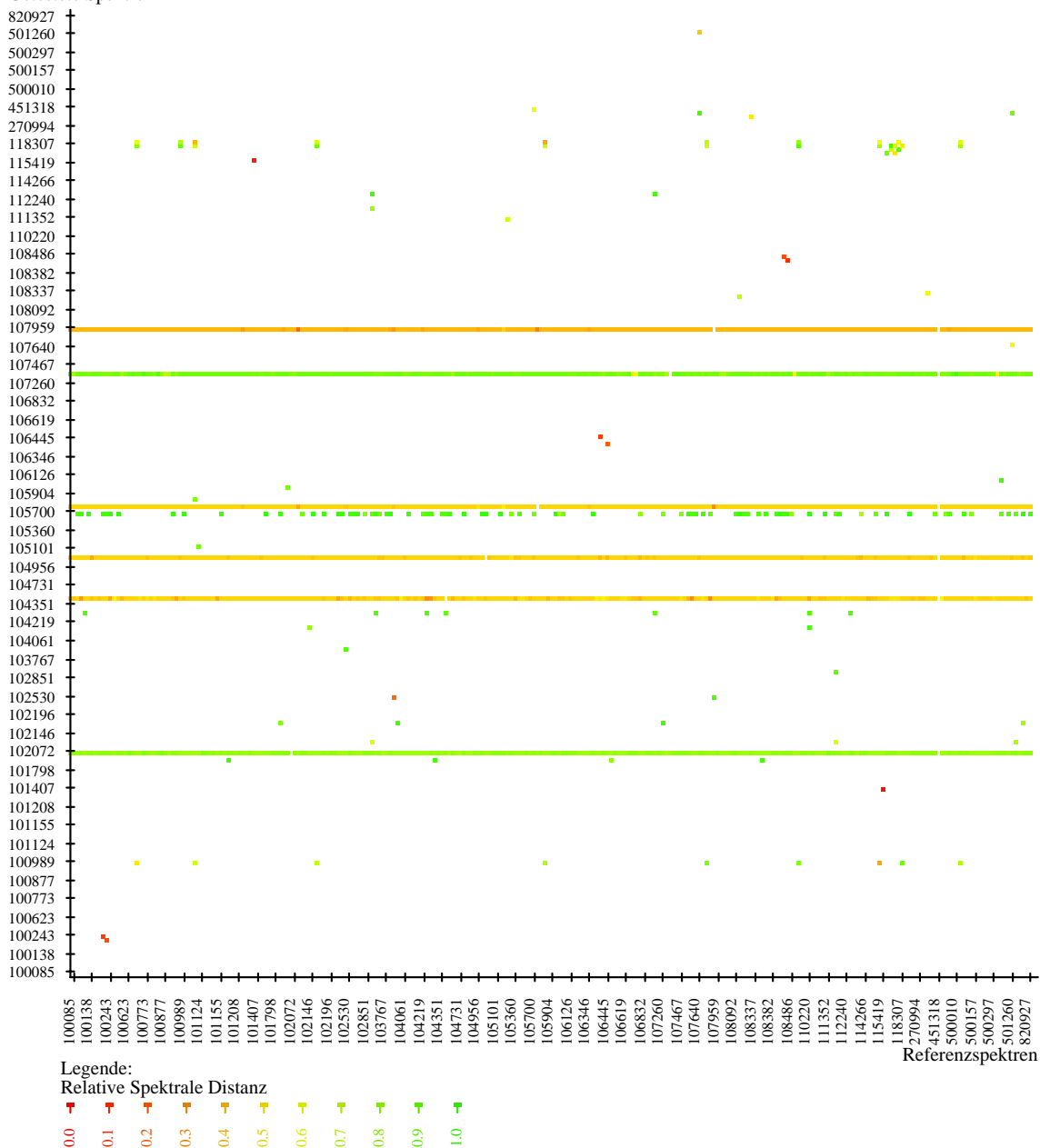
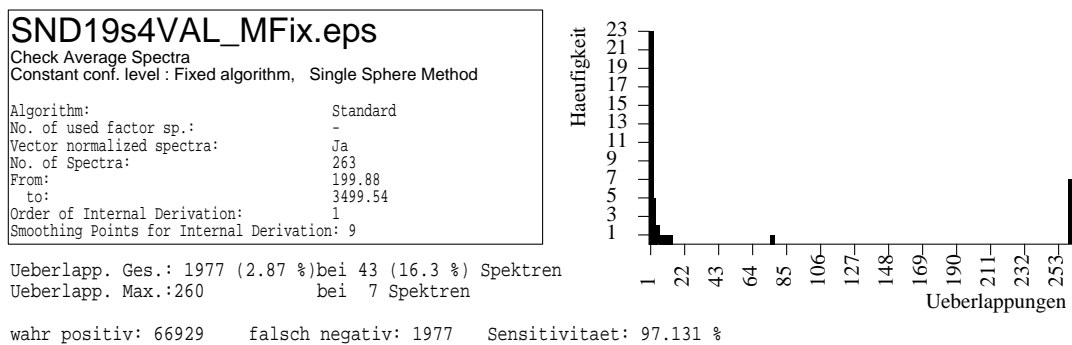


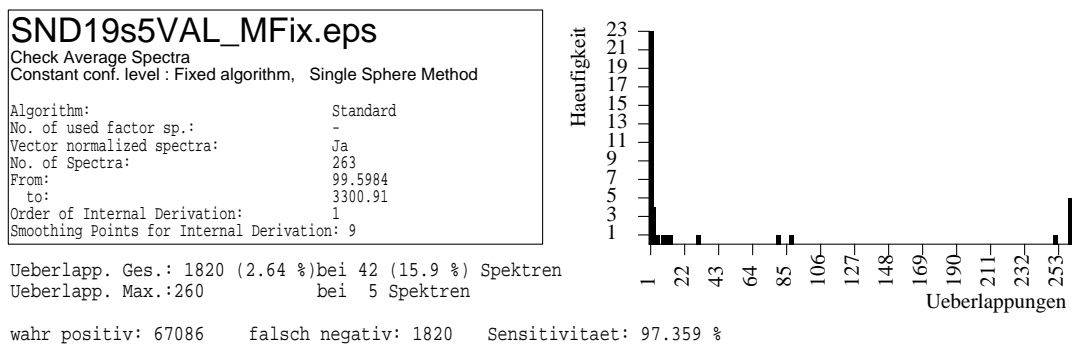
Abb. R23: Standardmethode (Vektornormierung, 1. Ableitung, 3500 cm^{-1} bis 150 cm^{-1}).



Getestete Spektren



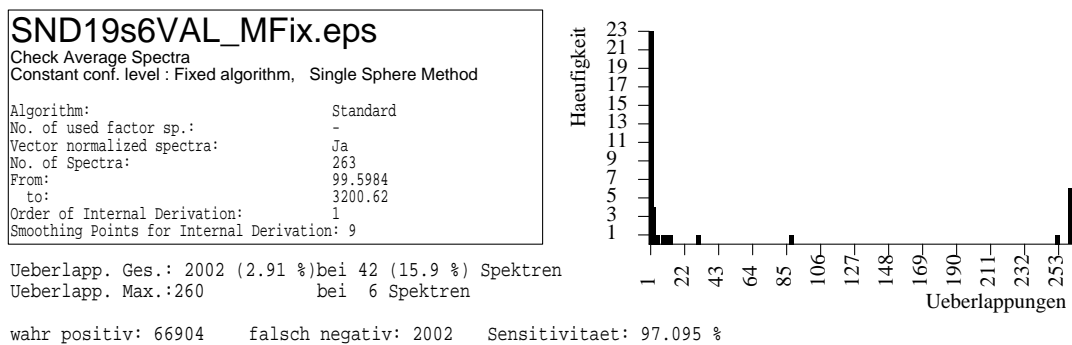
Abb. R24: Standardmethode (Vektornormierung, 1. Ableitung, 3500 cm^{-1} bis 200 cm^{-1}).



Getestete Spektren



Abb. R25: Standardmethode (Vektornormierung, 1. Ableitung, 3300 cm^{-1} bis 100 cm^{-1}).



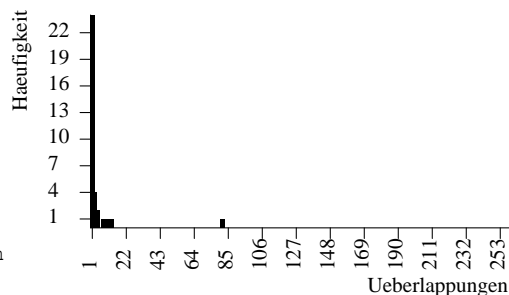
Getestete Spektren



Abb. R26: Standardmethode (Vektornormierung, 1. Ableitung, 3200 cm^{-1} bis 100 cm^{-1})

SND19s7VAL_MFix.eps
 Check Average Spectra
 Constant conf. level : Fixed algorithm, Single Sphere Method

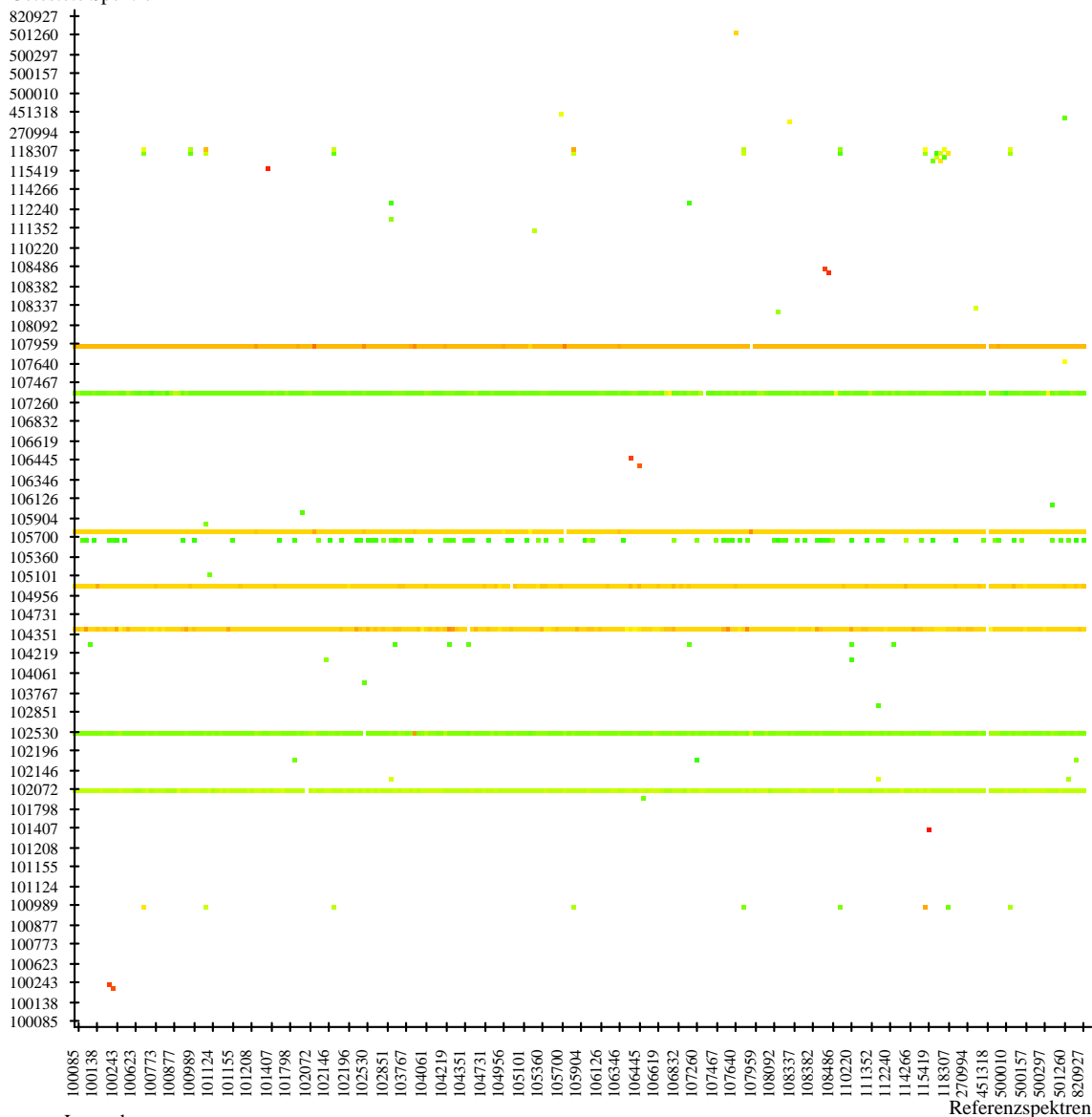
Algorithm: Standard
 No. of used factor sp.: -
 Vector normalized spectra: Ja
 No. of Spectra: 263
 From: 149.739
 to: 3200.62
 Order of Internal Derivation: 1
 Smoothing Points for Internal Derivation: 9



Ueberlapp. Ges.: 1977 (2.87 %) bei 42 (15.9 %) Spektren
 Ueberlapp. Max.: 260 bei 7 Spektren

wahr positiv: 66929 falsch negativ: 1977 Sensitivitaet: 97.131 %

Getestete Spektren



Legende:
 Relative Spektrale Distanz

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Abb. R27: Standardmethode (Vektornormierung, 1. Ableitung, 3200 cm^{-1} bis 150 cm^{-1})

7.4.8 Raman-Binärstrings

2_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 1700, Min: 1 Max:144

Gepruefte Spektren

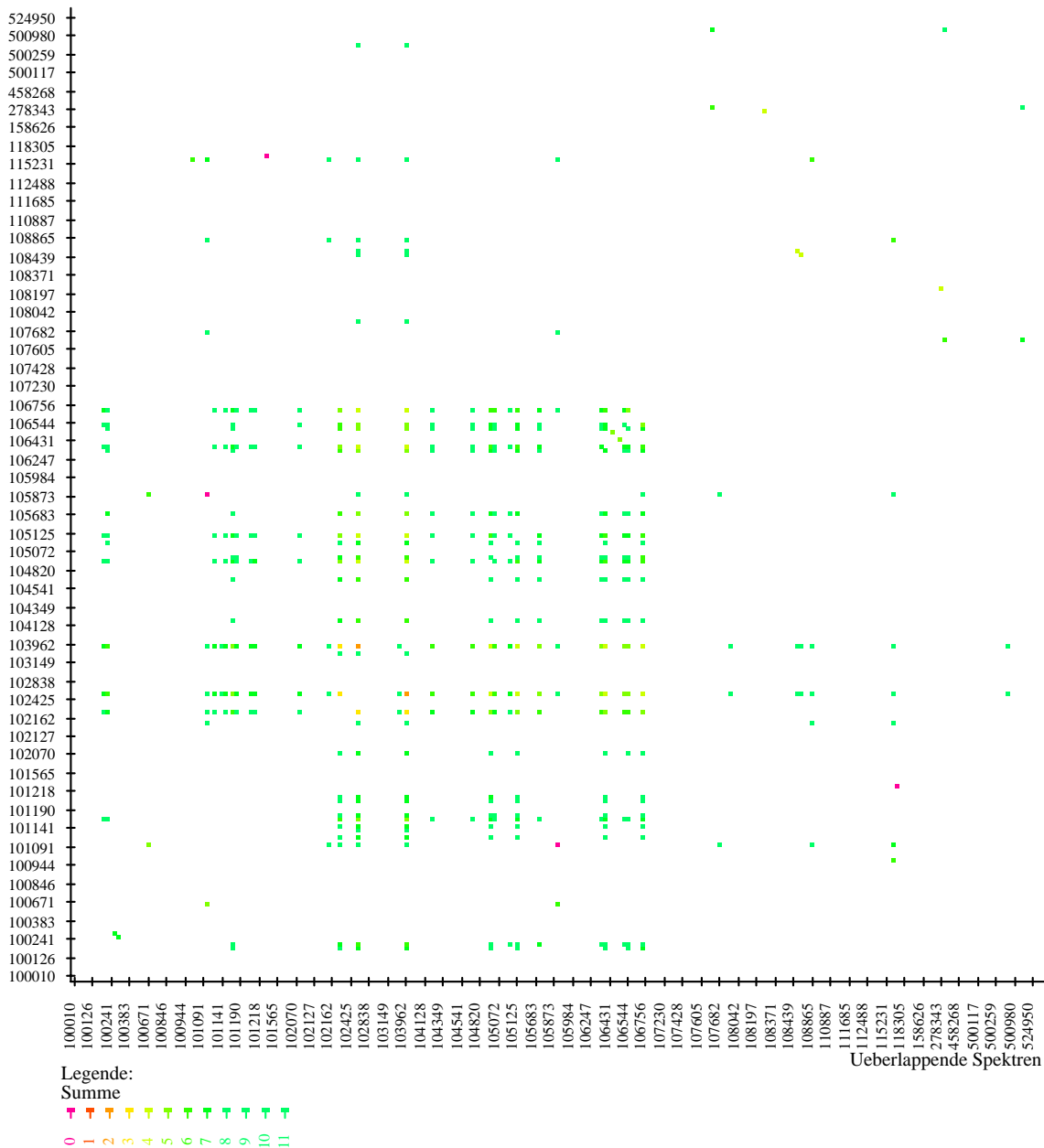


Abb. B1: Raman-Binärstring (2 cm^{-1}).

5_XOR.eps

O. Mandal
 ra_comp Version Jan 15 1999 11:10:55
 Intervallzahl: 682, Min: 1 Max:109

Gepruefte Spektren

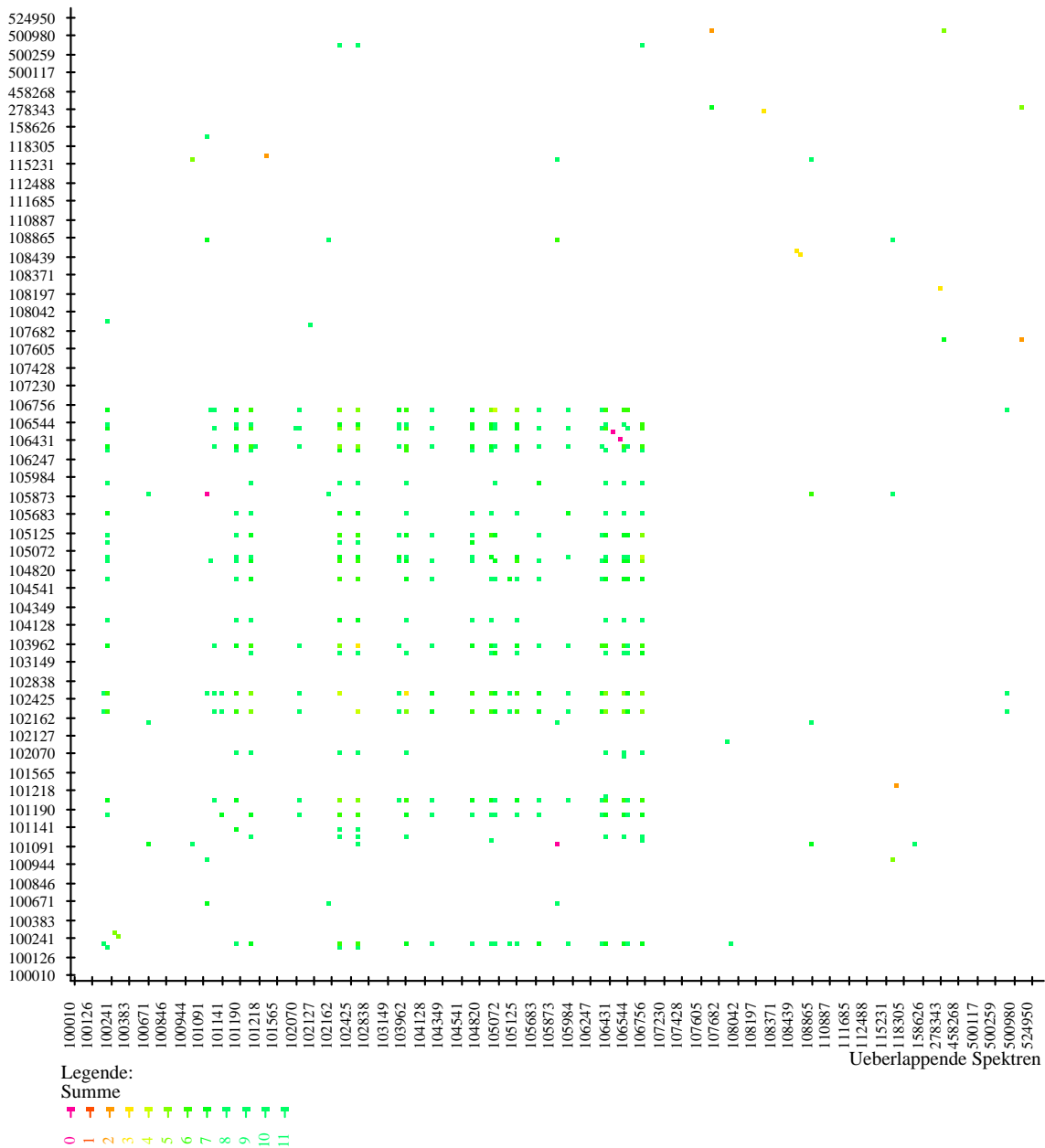


Abb. B2: NIR/Raman-Binärstring (5 cm^{-1}).

20_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 170, Min: 1 Max:102

Gepruefte Spektren

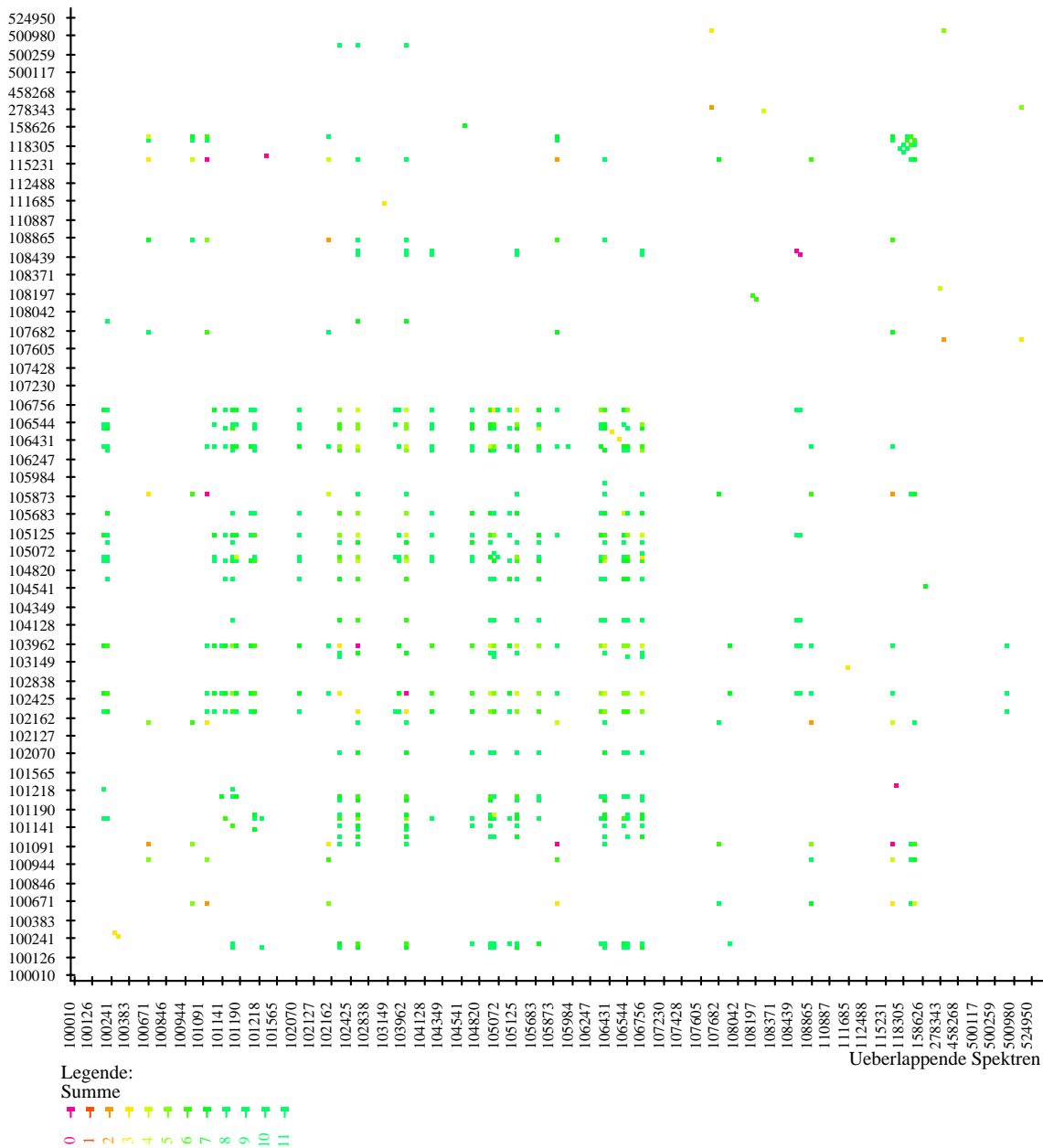


Abb. B3: Raman-Binärstring (20 cm^{-1}).

50_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 68, Min: 1 Max: 59

Gepruefte Spektren

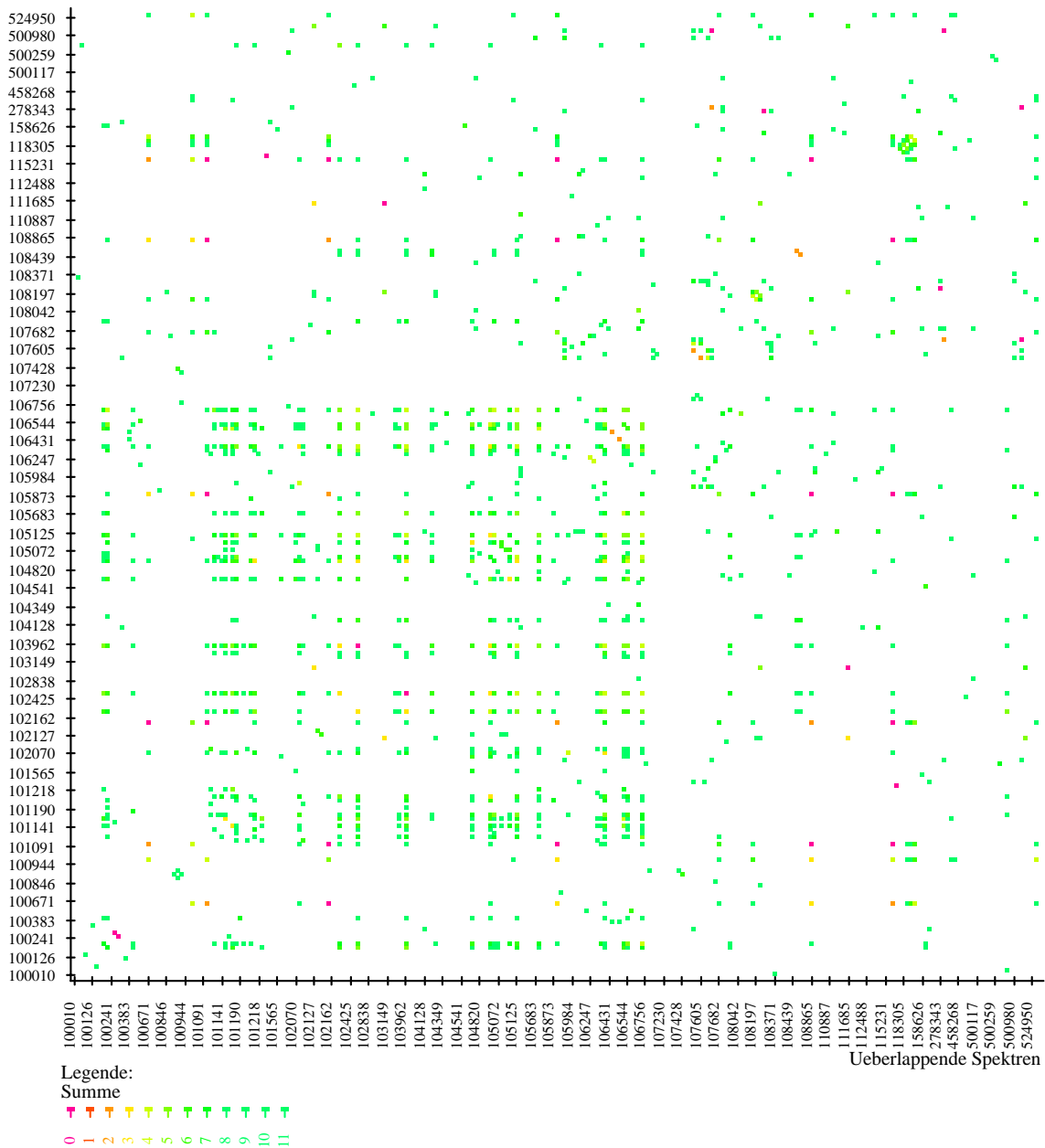


Abb. B4: Raman-Binärstring (50 cm^{-1}).

7.4.9 NIR-Binärstrings

2_XOR.eps

O. Mandal
ra_comp Version Jan 14 1999 15:35:21
Intervallzahl: 3999, Min: 1 Max: 54

Gepruefte Spektren

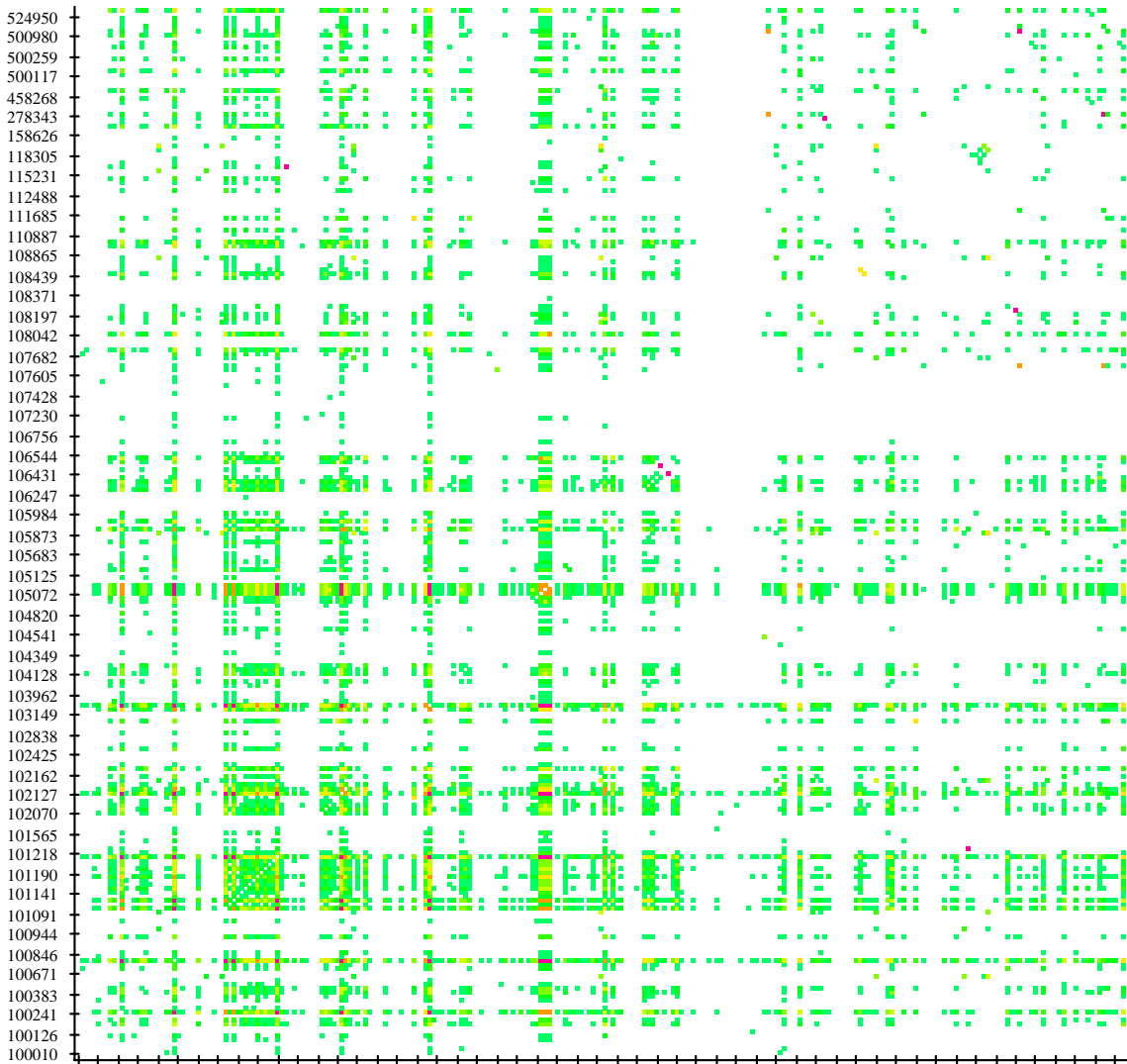


Abb. B5: NIR-Binärstring (2 cm^{-1}).

20_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 399, Min: 1 Max: 50

Gepruefte Spektren



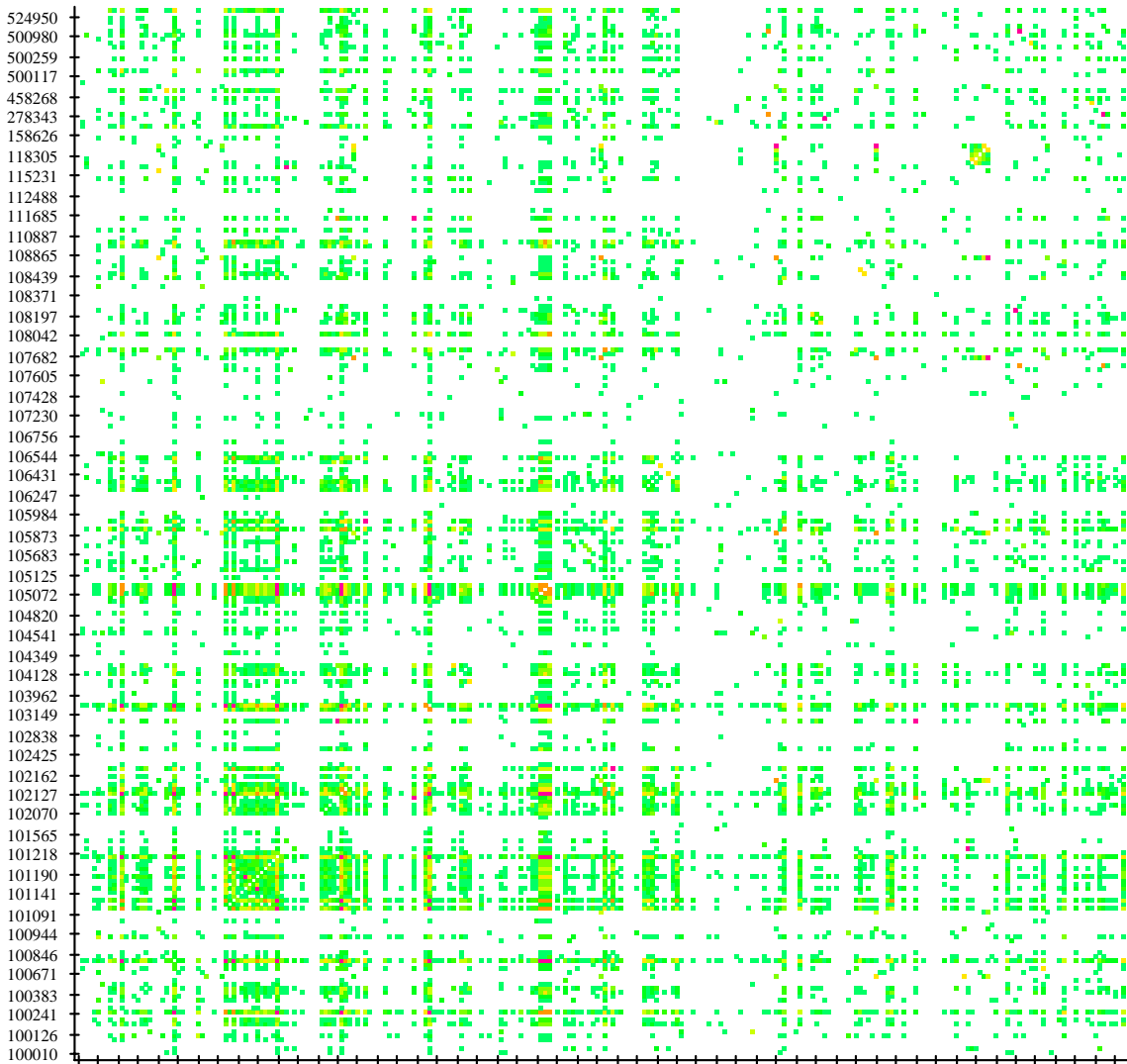
Legende:
 Summe
 0 1 2 3 4 5 6 7 8 9 10 11

Abb. B6: NIR-Binärstring (20 cm⁻¹).

50_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 159, Min: 1 Max: 39

Gepruefte Spektren



Legende:
 Summe
 0 1 2 3 4 5 6 7 8 9 10 11

Abb. B7: NIR-Binärstring (50 cm^{-1}).

7.4.10 Kombinierte NIR/Raman-Binärstrings

2_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 5700, Min: 3 Max:164

Gepruefte Spektren

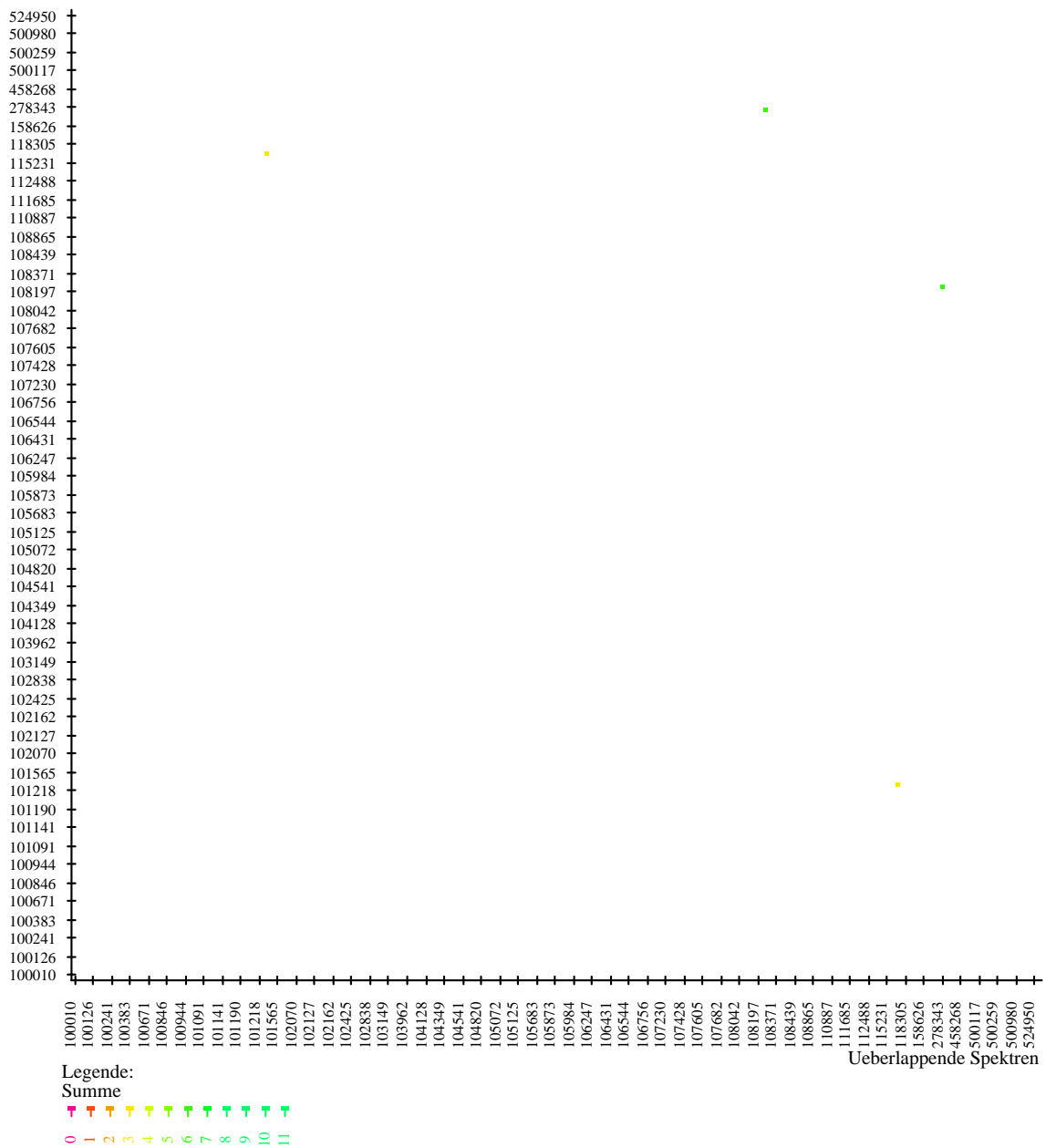


Abb. B8: NIR/Raman-Binärstring (2 cm^{-1}).

20_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 570, Min: 1 Max:128

Gepruefte Spektren



Abb. B9: NIR/Raman-Binärstring (20 cm^{-1}).

50_XOR.eps

O. Mandal
 ra_comp Version Jan 14 1999 15:35:21
 Intervallzahl: 228, Min: 1 Max: 81

Gepruefte Spektren

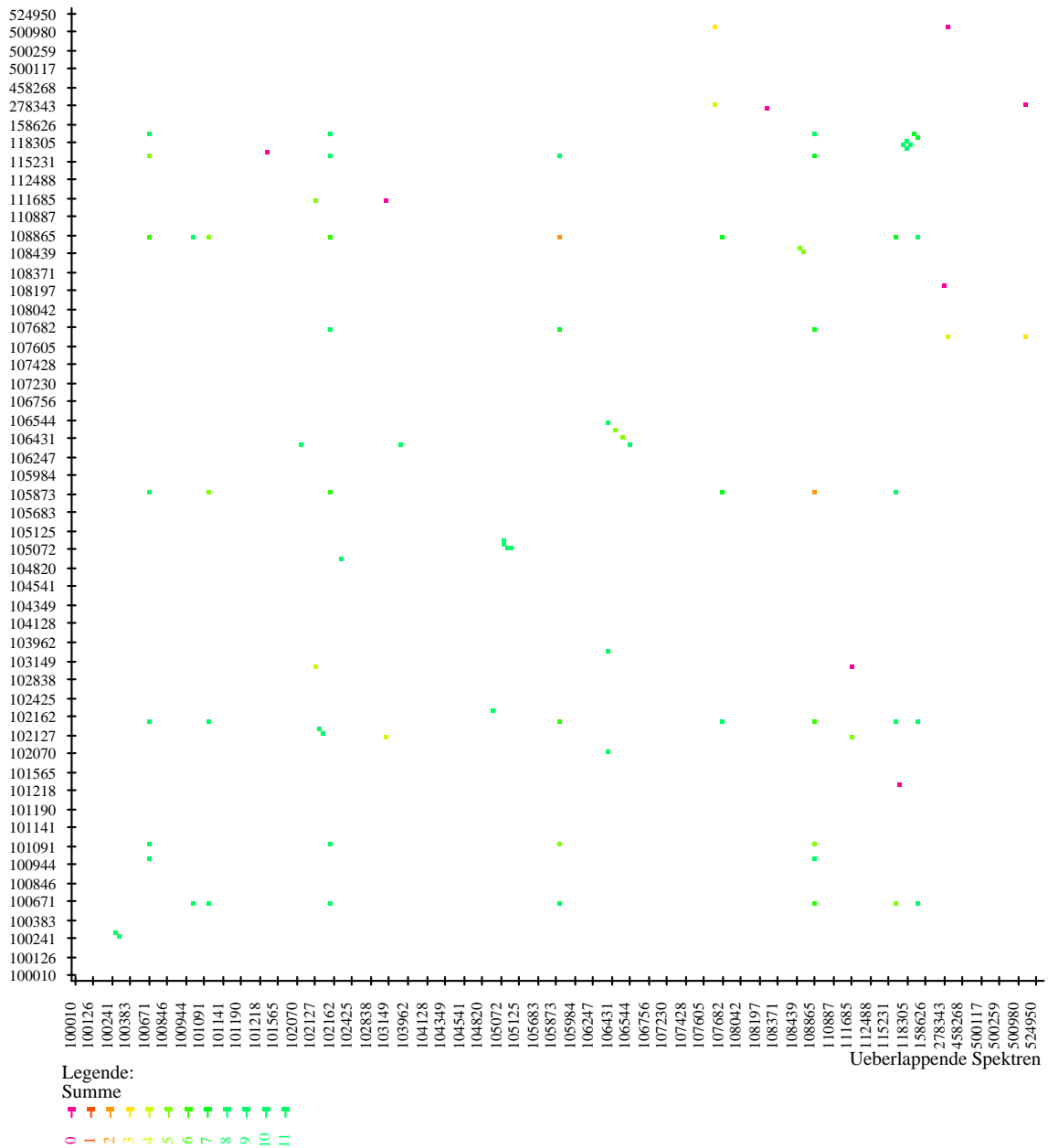


Abb. B10: NIR/Raman-Binärstring (50 cm^{-1}).