## Supplemental files submitted with Knaup et al.

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| family \# | no. of <br> affected <br> individuals <br> (SNaPshot <br> confirmed) | no. of <br> affected <br> generations | haplotype <br> analyses |
| :--- | :--- | :--- | :--- |
| A-29 (1*) | $>50(27)$ | 6 | yes |
| A-30 (2*) | $9(7)$ | 4 | yes |
| A-49 (5*) | $13(4)$ | 4 |  |
| A-33 (10*) | $7(2)$ | 3 |  |
| A-05 | $2(2)$ | 3 |  |
| A-11 | $3(3)$ | 3 | yes |
| A-12 | $4(1)$ | 3 |  |
| A-14 | $4(2)$ | 3 |  |
| A-21 | $5(2)$ | 4 |  |
| A-23 | $4(1)$ | 3 |  |
| A-24 | $3(1)$ | 2 |  |
| A-36 | $2(1)$ | 2 |  |
| A-37 | $>10(3)$ | 2 |  |
| A-38 | $4(2)$ | $>2$ |  |
| A-43 | $17(9)$ | 3 |  |
| A-47 | $7(4)$ | 3 |  |
| A-50 |  | 4 |  |

## Supplementary Table 1: List of ADTKD-MUC1 families

17 families with at least two affected successive generations were evaluated for ADTKD-MUC1 via SNaPshot minisequencing. Numbers provided are in many cases derived from patient history taking, where the affected status appeared very likely. Number of affected family members confirmed by SNaPshot minisequencing are given in parenthesis. In selected families haplotype reconstruction was performed by microsatellite analysis. *indicates the nomenclature of renamed families as cited in (Ekici et al. ${ }^{10}$ ).

Supplementary Table 2: List of antibodies (Ab) applied in our study

|  | company (clone) | host | dilution IB | dilution IHC/IF |
| :---: | :---: | :---: | :---: | :---: |
| Primary Ab: |  |  |  |  |
| 11ß HSD | Millipore (AB1296), Darmstadt, Germany | sheep |  | 1:200 |
| alpha Tubulin | GeneTex (GT114), Irvine, USA | mouse | 1:10000 |  |
| Megalin | Acris (DM3613P), Herford, Germany | mouse |  | 1:5000 |
| Aquaporin-2 | Sigma (SAB5200110), Darmstadt, Germany | rabbit |  | 1:1000 |
| MUC1 WT (VNTR) | Cell Signaling (VU4H5), Danvers, USA | mouse | 1:1000 | 1:100/1:100 |
| MUC1 WT (C-term) | Abcam (ab80952), Cambridge, UK | Armenian hamster |  | 1:1000/1:100 |
| MUC1-fs (pAb1-4) | n.a. | rabbit | 1:5000 | 1:5000/1:1000 |
| THP/UMOD | MP Cappel (55140), Cambridge, UK | goat |  | 1:1000/1:50 |
| $\beta$-Actin | Sigma (clone AC-15, 5441), Darmstadt, Germany | mouse | 1:10000 |  |
| Secondary Ab: |  | target/host |  |  |
| swine Anti-rabbit HRP | Dako (P0447), Santa Clara, USA | rabbit/swine | 1:2000 |  |
| goat Anti-mouse HRP | Dako (P0448), Santa Clara, USA | mouse/goat | 1:2000 |  |
|  |  |  |  |  |
| rb Anti-sheep IgG biotinylated | VectorLabs (BA-6000), Burlingame, USA | sheep/rabbit |  | 1:500 |
| rb Anti-goat biotinylated | VectorLabs (BA-5000), Burlingame, USA | goat/rabbit |  | 1:50 |
| Armenian hamster biotinylated | Jackson ImmunoResearch (127-065-160), West Grove, USA | armenian hamster/goat |  | 1:1000 |
| Alexa Fluor 594, Molecular Probes | Thermo Fisher Scientific (A28175), Darmstadt, Germany | mouse/goat |  | 1:500 |
| Alexa Fluor 488, Molecular Probes | Thermo Fisher Scientific (A11055) ), Darmstadt, Germany | goat/donkey |  | 1:500 |
| Alexa Fluor 488, Molecular Probes | Thermo Fisher Scientific (A11070) ), Darmstadt, Germany | rabbit/goat |  | 1:500 |
| Alexa Fluor 647 | Abcam (ab173004), Cambridge, UK | Armenian hamster/goat |  | 1:200 |

Dilutions for immunoblotting (IB), Immunohistochemistry (IHC) and Immunofluorescence (IF) as indicated.

Supplementary Table 3: List of microsatellites used in haplotype reconstruction and their respective primer sequences

| $\#$ | Microsatellit <br> e ID | Genetic Position <br> $(\mathrm{cM})$ | Physical Position <br> (bp in hg19) | Size (bp) | Forward Primer (5'-3') | ReversePrimer (5'-3') |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| 1 | ATA42G12 | 139.02 | $107,075,628$ | $178-196$ | AGCTAGGCACTTGTTGATGG | ATGTTGGTCCACATGTACCC |
| 2 | D1S534 | 151.88 | $119,678,203$ | $196-216$ | AGCACATAGCAGGCACTAGC | CGATTGTGCCACTACACAGT |
| 3 | D1S305 | 159.32 | $154,281,903$ | $156-176$ | CCAGNCTCGGTATGTTTTTACTA | CTGAAACCTCTGTCCAAGCC |
| 4 | D1S2714 | 161.05 | $155,115,519$ | $172-182$ | ATGAATTGCTTGAGCCCA | AGCTATCCTCCCACCTCAGA |
| 5 | D1S1153 | 161.05 | $155,268,703$ | $270-328$ | CAGACGAGACCCTAGAGAG | GGATTATAGGCAAGAGCCAC |
| 6 | D1S2777 | 161.05 | $155,460,360$ | $224-274$ | GCACCACGGAACTCCAGTAT | CACCACTGTGCCCAGCTAAT |
| 7 | D1S303 | 161.05 | $155,637,498$ | $181-191$ | CGACAAGAGCGAAACTCCAT | GCTTCCCAGAGGCTAGGATT |
| 8 | D1S1595 | 161.05 | $155,688,364$ | $265-293$ | ATGGTATGAACCTGGAGGTG | GGCAGATAAAAGGACTGCAA |
| 9 | D1S2721 | 161.05 | $156,153,449$ | $201-247$ | TTGCTCGGCCAGAGTCT | ACGCATCACACCTGGCTAGT |
| 10 | D1S1679 | 170.84 | $162,367,764$ | $144-172$ | GCCATCAAGAAAACTAGTACTGC | ACCATGGTACTCAGCAGTGC |
| 11 | D1S1677 | 175.62 | $163,559,700$ | $184-212$ | AGTCAGCTTGATTGACCCAG | CTTAGTGTGACAGGAAGGACG |
| 12 | D1S1589 | 192.05 | $174,261,084$ | $199-217$ | TACTCAGGAGGCAGAGATGG | CTGCTTTGGGTTTCACTTGT |

Microsatellites 1-4 are proximal and 5-12 are distal of MUC1.

## Supplementary Table 4: List of microsatellites based genotypes

| ID | $\begin{aligned} & \text { ATA42G } \\ & 12 \end{aligned}$ | D1S534 | D1S305 | D1S2714 | D1S1153 | D1S2777 | D1S303 | D1S1595 | D1S2721 | D1S1679 | D1S1677 | D1S1589 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| A-11_II-2 | 184 | 208 | 172 | 180 | 324 | 266 | 183 | 290 | 232 | 148 | 208 | 200 |
|  | 187 | 202 | 172 | 180 | 315 | 266 | 187 | 282 | 242 | 156 | 212 | 210 |
| A-11_III-1 | 184 | 212 | 168 | 180 | 324 | 266 | 183 | 274 | 242 | 164 | 192 | 201 |
|  | 187 | 202 | 172 | 180 | 0 | 266 | 187 | 282 | 242 | 155 | 212 | 210 |
| A-11_III-2 | 0 | 0 | 172 | 180 | 297 | 268 | 187 | 282 | 236 | 164 | 200 | 200 |
|  | 184 | 0 | 172 | 180 | 324 | 266 | 183 | 290 | 232 | 148 | 208 | 200 |
| A-11_III-3 | 181 | 214 | 172 | 180 | 297 | 268 | 187 | 282 | 236 | 164 | 200 | 200 |
|  | 184 | 208 | 172 | 180 | 324 | 266 | 183 | 290 | 232 | 148 | 208 | 200 |
| A-11_III-4 | 184 | 212 | 168 | 180 | 324 | 266 | 183 | 274 | 0 | 164 | 0 | 200 |
|  | 187 | 0 | 172 | 180 | 0 | 266 | 187 | 282 | 242 | 156 | 212 | 210 |
| A-11_III-5 | 181 | 214 | 170 | 178 | 294 | 266 | 183 | 282 | 238 | 164 | 200 | 204 |
|  | 184 | 208 | 172 | 180 | 297 | 268 | 187 | 282 | 242 | 168 | 200 | 212 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 6 \end{aligned}$ | 184 | 212 | 160 | 174 | 312 | 272 | 181 | 276 | 246 | 160 | 200 | 208 |
|  | 187 | 204 | 162 | 180 | 297 | 268 | 181 | 282 | 234 | 148 | 212 | 200 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 7 \end{aligned}$ | 175 | 206 | 168 | 146 | 294 | 266 | 181 | 276 | 242 | 156 | 196 | 200 |
|  | 187 | 204 | 162 | 142 | 318 | 268 | 189 | 286 | 240 | 168 | 204 | 216 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 8 \end{aligned}$ | 175 | 206 | 168 | 146 | 0 | 266 | 181 | 276 | 242 | 156 | 196 | 200 |
|  | 184 | 200 | 172 | 142 | 0 | 268 | 189 | 286 | 234 | 168 | 200 | 200 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 9 \end{aligned}$ | 181 | 212 | 164 | 178 | 321 | 260 | 185 | 264 | 234 | 160 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 11 \end{aligned}$ | 181 | 212 | 164 | 178 | 321 | 260 | 185 | 264 | 234 | 160 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 13 \end{aligned}$ | 181 | 216 | 174 | 172 | 312 | 268 | 181 | 282 | 240 | 160 | 204 | 200 |
|  | 184 | 200 | 162 | 180 | 318 | 268 | 185 | 282 | 234 | 172 | 200 | 210 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 15 \end{aligned}$ | 184 | 216 | 162 | 178 | 324 | 266 | 185 | 274 | 234 | 148 | 196 | 200 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |


| $\begin{aligned} & \text { A-29_IV- } \\ & 17 \end{aligned}$ | 184 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 148 | 192 | 200 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 187 | 206 | 172 | 182 | 315 | 266 | 185 | 286 | 234 | 168 | 196 | 210 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 18 \end{aligned}$ | 0 | 0 | 168 | 178 | 312 | 266 | 185 | 286 | 228 | 148 | 192 | 200 |
|  | 187 | 206 | 172 | 182 | 315 | 266 | 185 | 286 | 234 | 168 | 196 | 210 |
| $\begin{aligned} & \text { A-29_IV- } \\ & 19 \end{aligned}$ | 181 | 204 | 168 | 178 | 312 | 266 | 185 | 286 | 228 | 0 | 192 | 200 |
|  | 187 | 206 | 172 | 182 | 315 | 266 | 185 | 286 | 234 | 168 | 196 | 210 |
| A-29_V-1 | 184 | 206 | 158 | 142 | 288 | 268 | 185 | 272 | 234 | 172 | 204 | 196 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| A-29_V-3 | 184 | 206 | 158 | 142 | 288 | 268 | 185 | 272 | 234 | 172 | 204 | 196 |
|  | 187 | 212 | 164 | 142 | 310 | 272 | 185 | 290 | 242 | 152 | 196 | 212 |
| A-29_V-4 | 184 | 206 | 158 | 142 | 288 | 268 | 185 | 272 | 234 | 172 | 204 | 196 |
|  | 187 | 212 | 164 | 142 | 310 | 272 | 185 | 290 | 242 | 152 | 196 | 212 |
| A-29_V-6 | 187 | 212 | 174 | 180 | 288 | 268 | 185 | 272 | 246 | 152 | 196 | 200 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| A-29_V-8 | 187 | 212 | 174 | 180 | 288 | 268 | 185 | 272 | 246 | 152 | 196 | 200 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| A-29_V-9 | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
|  | 187 | 212 | 174 | 180 | 288 | 268 | 185 | 272 | 246 | 152 | 196 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 11 \end{aligned}$ | 189 | 208 | 162 | 180 | 312 | 266 | 185 | 288 | 240 | 160 | 196 | 212 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 13 \end{aligned}$ | 187 | 204 | 162 | 180 | 297 | 268 | 181 | 282 | 234 | 148 | 212 | 200 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 14 \end{aligned}$ | 184 | 212 | 160 | 174 | 312 | 272 | 181 | 276 | 246 | 160 | 200 | 208 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 15 \end{aligned}$ | 184 | 200 | 162 | 180 | 318 | 266 | 193 | 278 | 232 | 156 | 204 | 200 |
|  | 181 | 212 | 164 | 178 | 321 | 260 | 185 | 264 | 234 | 160 | 204 | 204 |
| $\begin{aligned} & \text { A-29_V- } \\ & 16 \end{aligned}$ | 181 | 212 | 164 | 178 | 321 | 260 | 185 | 264 | 234 | 160 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 17 \end{aligned}$ | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
|  | 189 | 213 | 163 | 178 | 303 | 269 | 185 | 284 | 240 | 156 | 204 | 204 |
| A-29_V- | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |


| 18 | 184 | 200 | 162 | 180 | 318 | 268 | 185 | 282 | 234 | 172 | 200 | 210 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\begin{aligned} & \text { A-29_V- } \\ & 19 \end{aligned}$ | 184 | 200 | 162 | 180 | 318 | 268 | 185 | 282 | 234 | 172 | 200 | 210 |
|  | 184 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_V- } \\ & 20 \end{aligned}$ | 184 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 148 | 192 | 200 |
|  | 187 | 206 | 166 | 178 | 314 | 266 | 193 | 284 | 232 | 148 | 196 | 210 |
| $\begin{aligned} & \text { A-29_V- } \\ & 21 \end{aligned}$ | 181 | 200 | 172 | 184 | 318 | 266 | 193 | 276 | 242 | 148 | 192 | 200 |
|  | 187 | 206 | 172 | 182 | 315 | 266 | 185 | 286 | 234 | 168 | 196 | 210 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 1 \end{aligned}$ | 184 | 212 | 169 | 180 | 321 | 262 | 189 | 284 | 234 | 176 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 204 | 196 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 2 \end{aligned}$ | 184 | 212 | 162 | 142 | 324 | 266 | 185 | 286 | 234 | 164 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 3 \end{aligned}$ | 184 | 212 | 162 | 142 | 324 | 266 | 185 | 286 | 234 | 164 | 204 | 204 |
|  | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 4 \end{aligned}$ | 184 | 206 | 158 | 142 | 288 | 268 | 185 | 172 | 234 | 172 | 204 | 196 |
|  | 186 | 210 | 172 | 180 | 312 | 272 | 191 | 288 | 242 | 156 | 200 | 204 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 5 \end{aligned}$ | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 152 | 200 | 200 |
|  | 187 | 203 | 173 | 180 | 293 | 267 | 185 | 296 | 234 | 160 | 192 | 200 |
| $\begin{aligned} & \mathrm{A}-29 \_\mathrm{VI}- \\ & 6 \end{aligned}$ | 184 | 206 | 164 | 180 | 318 | 266 | 197 | 284 | 242 | 156 | 192 | 212 |
|  | 187 | 212 | 174 | 180 | 288 | 268 | 185 | 272 | 246 | 152 | 196 | 200 |
| $\begin{aligned} & \hline \text { A-29_VI- } \\ & 7 \end{aligned}$ | 187 | 212 | 174 | 180 | 288 | 268 | 185 | 272 | 246 | 152 | 196 | 200 |
|  | 184 | 206 | 164 | 180 | 318 | 266 | 197 | 284 | 242 | 156 | 192 | 212 |
| $\begin{aligned} & \text { A-29_VI- } \\ & 8 \end{aligned}$ | 187 | 200 | 172 | 174 | 318 | 266 | 197 | 276 | 242 | 156 | 196 | 200 |
|  | 187 | 213 | 175 | 178 | 324 | 266 | 193 | 288 | 244 | 160 | 204 | 200 |
| A-30_III-1 | 181 | 202 | 174 | 178 | 318 | 266 | 192 | 274 | 244 | 155 | 200 | 210 |
|  | 184 | 206 | 164 | 176 | 318 | 268 | 184 | 286 | 238 | 143 | 208 | 196 |
| A-30_III-2 | 184 | 206 | 158 | 178 | 0 | 268 | 184 | 278 | 234 | 155 | 196 | 200 |
|  | 187 | 200 | 168 | 178 | 330 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| A-30_III-4 | 184 | 200 | 172 | 178 | 309 | 266 | 184 | 290 | 236 | 155 | 196 | 200 |
|  | 187 | 200 | 168 | 178 | 330 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| A-30_III-5 | 184 | 198 | 172 | 176 | 312 | 266 | 184 | 282 | 242 | 151 | 200 | 200 |
|  | 181 | 198 | 164 | 180 | 297 | 268 | 180 | 290 | 238 | 155 | 200 | 204 |


| A-30_III-6 | 184 | 200 | 172 | 178 | 0 | 266 | 184 | 290 | 236 | 147 | 196 | 200 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 187 | 200 | 168 | 178 | 0 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| A-30_III-7 | 184 | 206 | 158 | 178 | 0 | 268 | 184 | 278 | 234 | 0 | 196 | 200 |
|  | 187 | 200 | 168 | 178 | 0 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| $\begin{aligned} & \text { A-30_IV- } \\ & 1 \end{aligned}$ | 184 | 206 | 164 | 176 | 318 | 268 | 184 | 286 | 238 | 143 | 208 | 196 |
|  | 187 | 200 | 168 | 178 | 0 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| $\begin{aligned} & \text { A-30_IV- } \\ & 2 \end{aligned}$ | 184 | 200 | 172 | 178 | 309 | 266 | 184 | 290 | 236 | 155 | 196 | 200 |
|  | 187 | 200 | 168 | 178 | 330 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| $\begin{aligned} & \text { A-30_IV- } \\ & 3 \end{aligned}$ | 184 | 198 | 172 | 176 | 312 | 266 | 184 | 282 | 242 | 151 | 0 | 200 |
|  | 187 | 200 | 168 | 178 | 0 | 266 | 184 | 290 | 242 | 167 | 200 | 200 |
| $\begin{aligned} & \text { A-30_IV- } \\ & 4 \end{aligned}$ | 184 | 206 | 158 | 178 | 320 | 268 | 184 | 0 | 234 | 0 | 0 | 200 |
|  | 181 | 208 | 162 | 180 | 326 | 262 | 180 | 0 | 233 | 167 | 204 | 204 |

All genotypes used in haplotype reconstruction for members of families A-11, A-29 and A-30. The first column show the merged family and individual IDs separated with an underscore as indicated in the pedigrees in figure 2. Columns 2-13 show the genotypes as allele fragment length (bp) of respective microsatellites.

## SUPPLEMENTARY METHODS

Cell culture and reagents. HeLa cells were purchased from the German Collection of Microorganisms and Cell Cultures (DSMZ, Braunschweig, Germany). The cells were cultivated in DMEM, 1.0 g glucose/L, $10 \%$ fetal calf serum, $2 \mathrm{mmol} / \mathrm{L} \mathrm{L}-$ glutamine, 100 U penicillin and $100 \mu \mathrm{~g}$ streptomycin $/ \mathrm{ml}$. Medium and penicillin/streptomycin were supplied by PAN-Biotech (Aidenbach, Germany), fetal calf serum (standard "Gold") by PAA Laboratories (Coelbe, Germany). If not stated otherwise, all reagents were purchased from Sigma-Aldrich (Taufkirchen, Germany). Human urinary primary tubular cells (hUPTs) were generated and cultivated as described by Zhou et al. ${ }^{33}$

Cloning of MUC1 cDNA's. The cDNA coding for the human MUC1 gene including 22 tandem repeats (MUC1/22TR) was a kind gift from Prof. Dr. Olivera Finn (University of Pittsburgh) and is described elsewhere ${ }^{34}$. For the generation of the MUC1/22TR-fs plasmids, a PCR-Fragment generated by the following primers: 5'TAA TAC GAC TCA CTA TAG GG $-3^{\prime}$ and $5^{\prime}$ - GG GGC TGA GGT TGA CAT CGT GGG CTG $-3^{\prime}$ was amplified using the MUC1/22TR cDNA as a template. Primer 1 binds to the T7 promoter within the backbone of pcDNA3, the second primer contains one additional basepair and covers the restriction site BbvCl in order to generate a frameshift directly upstream of the VNTR. The amplified PCR fragment was digested with the restriction enzymes HindIII and BbvCl and cloned into the original MUC1/22TR cDNA. An internal Flag-Tag (DYKDDDDK) followed by the restriction site Sacll was fused directly to the end of exon 2 in analog to the method of Burdick and colleagues ${ }^{35}$ into the MUC1/22/TR-wt and -fs plasmid by PCR-mutagenesis using the following primers: $5^{\prime}$ - TAC ATC AAT GGG CGT GGA TA -3', $5^{\prime}$ - A GGC CGG GGC TGG CTT GTT GTC - $3^{\prime}, 5^{\prime}$ - CTT GTC GTC GTC ATC CTT GTA ATC AGC ATT CTT CTC AGT AGA -3' and $5^{\prime}$ - GAT GAC GAC GAC AAG CCG CGG GCT GTG AGT ATG ACC AGC AGC -3' and the restriction sites HindIII and BbvCl. For cloning, PCR fragments were purified using NucleoSpin Extract II from Macherey-Nagel (Düren, Germany). All plasmids were sequenced by standard procedueres at GATC Biotech (Konstanz, Germany).

Generation of antibodies. Four different rabbits were immunized with a synthetic peptide against the following amino acid sequence: $\mathrm{NH}_{2}{ }^{-}$ CHLGPGHQAGPGLHRPPSPR-CONH 2 , corresponding to the VNTR of the frameshift MUC1 protein (MUC1-fs). All animals were immunized at day 1, 60 and 90. For testing, serum was taken 60, 90 or 120 days after first injection. To obtain MUC1-fs antibodies, all sera of day 120 were affinity purified with a column coupled with the synthetic MUC1-fs VNTR peptide. All steps were carried out commercially by Pineda-antibody-service (Pineda, Berlin, Germany).

Human samples. Kidney biopsies were collected retrospectively. The study was approved by the local ethics committee (protocol no. 4103 and $181 \_15 \mathrm{Bc}$ ).

Transfection and Immunoblotting. HeLa cells were transfected either with equal amounts of empty vector (pcDNA3), wildtype human MUC1 cDNA (MUC1/22TR, 22 repeats of VNTR) or frameshift human MUC1 cDNA (MUC1/22TR-FS) using jetPEI transfection reagent (Polyplus-transfection, Illkirch, France) according to the manufacturer's instruction. 24 hours after transfection, cells were washed twice with PBS and homogenized into extraction buffer (8 M urea, 10\% glycerol, 1\% SDS, 10 mM TrisHCl pH 6.8, protease inhibitor complete ${ }^{\mathrm{TM}}$ (Roche, Mannheim, Germany)). Protein concentration was measured according to the manufacturer's manual using the DC Protein Assay (BioRad, California, USA). Equal amounts of protein were separated by SDS PAGE, transferred to PVDF membranes (Millipore, Bedford, MA, USA) and stained with antibodies against the wildtype MUC1 VNTR (MUC1 (VU4H5) mAb; no. 4538; Cell Signaling, Danvers, MA, USA; dilution 1:1000) or one of four antibodies (dilution 1:5000) generated to detect the MUC1-fs protein (pAb1-4-fs). IVTT extracts were generated following manufacturers description (TNT, T7 Quick Coupled Transcription/Translation System, Promega, Wisconsin, USA) using linearized plasmids for MUC1/22TR or MUC1/22TR-FS). For antibody testing, pure serum of immunized rabbits was used 1:20.000. As loading control, $\beta$-actin was stained using the monoclonal antibody $\beta$-actin (dilution: 1:10.000, Sigma Aldrich, St. Louis, USA no. A5441). Signals were visualized by the ECL system from GE Healthcare (Munich, Germany). For detailed antibody information, see supplementary table 2.
siRNA knockdown in hUPTs (human urinary primary tubular cells). hUPTs were seeded 24 hours prior to transfection for siRNA experiments and knockdown was performed as reported previously by Warnecke et al. ${ }^{36}$. Following siRNAs were used: siMUC1_5 Cat. No.: SI00162988, siMUC1_7 Cat. No: SI02780673, siMUC1_10 Cat. No.: SI04949826 and siMUC1_11 Cat. No.: SI04949833 (all Qiagen, Hilden, Germany).

Immunohistochemistry. Paraffin sections ( $2 \mu \mathrm{~m}$ ) were dewaxed in xylene and rehydrated in a series of ethanol washes. For antigen retrieval before staining of MUC1-wt VU4H5 and -fs protein, all slides were cooked in a microwave 20 min in 0.1 M citrate puffer. For antigen retrieval before staining MUC1-wt ab80952, THP and 11 BHSD, slides were cooked for 7 or 5 min in $1 \times$ TRS (Target Retrieval Solution, DAKO, Glostrup, Denmark) in a standard pressure cooker. In the case of MUC1-wt ab80952 and THP staining, an additional avidin and biotin blocking was performed (Biotin Blocking System, DAKO). Endogenous peroxidase activity was blocked by incubation for 10 min with Peroxidase Real (DAKO). Slides for the MUC1-wt VU4H5 and -fs protein where blocked with $2.5 \%$ Normal Horse Serum (Vector Laboratories) for 20 min , for 11ßHSD 10\% Normal Rabbit Serum was used for 30 min and for Muc1-wt ab80952 as well as THP Protein Block (DAKO) was use. All blocking steps where performed at room temperature.
Sections were incubated with primary antibodies (see Supplementary Table 2) diluted in Antibody Diluent (DAKO) overnight at $4^{\circ} \mathrm{C}$ (MUC1-wt VU4H5, ab80952, MUC1-fs, THP) or at $37^{\circ} \mathrm{C}$ for 60 min (11ßHSD). In case of THP, 11ßHSD and MUC1-wt ab80952, sections were incubated with biotinylated secondary antibody for 30min at room temperature. Next, slides were incubated with streptavidin/biotinylated alkaline phosphatase (Vectastain Elite ABC Kit, Vector Laboratories, Burlingame, USA) for 30 min . For MUC1-wt VU4H5 and MUC1-fs protein staining, slides were incubated with ImmPRESS Reagent Kit Anti-mouse, -rabbit Ig for 30 min. Finally, AEC+ solution (DAKO) or DAB solution (ImmPACT DAB Peroxidase Substrate Kit, Vector Laboratories) was used as chromogen according to the manufacturer's instructions. For identification of proximal tubules and collecting ducts sections of paraffin-embedded renal biopsies were immunohistochemically stained using antibodies specific for Megalin (mouse monoclonal, Acris, DM3613P, 1:5000) and Aquaporin-2 (rabbit polyclonal, Sigma, SAB5200110, 1:1000). All biopsies were fixed
and stained according to our standard techniques (Ventana BenchMark ULTRA stainer and UltraView DAB IHC Detection Kit, Roche).
All incubations were performed in a humidified chamber. Between incubations, specimens were washed three times in Tris-buffered saline ( $50 \mathrm{mmol} / \mathrm{L}$ Tris- HCl and $136 \mathrm{mmol} / \mathrm{L} \mathrm{NaCl}, \mathrm{pH} 7.4$ ). Samples were processed in parallel throughout. Finally, the sections were counterstained with hematoxylin solution according to Mayer (DAKO) and analyzed with a Leica DMRB microscope (Leica, Bensheim, Germany).

Immunofluorescence. Paraffin sections ( $2 \mu \mathrm{~m}$ ) were dewaxed in xylene and rehydrated in a series of ethanol washes. Antigen retrieval was performed by cooking all slides for 10 min (MUC1-wt (VU4H5) and THP) or 20 min (MUC1-fs, MUC1-wt (ab80952) and THP) in 0.1 M citrate puffer using a microwave. Next all slides where blocked with sterile 1\% BSA/ PBS. Primary antibodies where diluted in 1\% BSA/ PBS and incubated overnight at $4^{\circ} \mathrm{C}$ (see Supplementary Table 2). Following the slides where incubated for 1.5 hours at room temperature in Alexa Fluor secondary antibodies diluted in 1\% BSA/PBS. For nuclear DNA counterstaining, DAPI (Thermo Fisher Scientific, D1306), diluted 1:1000 in PBS, was added for 4 min.
Samples were always processed in parallel and incubated in a humidified chamber. Between incubations, specimens were washed three times in PBS. A Leica DMRB microscope (Leica, Bensheim, Germany) was used for analyzing the slides.

Antibody pre-adsorption assay. Prior to routine immunohistochemistry (see above) the pAb3-fs was incubated on a roller platform with the MUC1-fs peptide $\left(\mathrm{NH}_{2}{ }^{-}\right.$ CHLGPGHQAGPGLHRPPSPR-CONH 2 ) used for generating the antibodies pAbx-fs, a hSPAG4 (human Sperm Associated Antigen 4) peptide $\left(\mathrm{NH}_{2}{ }^{-}\right.$ RSAEPGPGEPEGRRARGPSC-CONH ${ }_{2}$ ) also consisting of 20 amino acids (used for the generation of hSPAG4 antibody, described in Knaup et al. ${ }^{22}$ ) and "no peptide" (antibody diluent, DAKO) for 2 h at room temperature. The peptides were applied at a final concentration of $20 \mu \mathrm{~g} / \mathrm{ml}$. Subsequently the antibodies were applied at identical dilutions, as described above.

## SUPPLEMENTARY FIGURE LEGENDS

## Supplementary Figure 1: Partial co-localization of mucin 1 and THP in kidney

 tubules. Immunofluorescent staining of healthy control kidney shows partial colocalization of mucin 1 (red), stained with the C-term mucin 1 antibody and THP (green, Tamm Horsfall Protein), an established kidney tubule marker of the TAL (thick ascending limb). Arrows indicate kidney tubules with shared expression of both markers.
## Supplementary Figure 2: Localization of mucin 1 in kidney tubule system.

 Immunohistochemical analyses of serial human kidney sections for mucin 1 and distinct markers of the kidney tubule system (megalin, THP (Tamm Horsfall Protein), 11ßHSD (11-ß-hydroxysteroid dehydrogenase) and aquaporin-2. Partial colocalization is visible for mucin 1 and THP (TAL, thick ascending limb) as well as for aquaporin-2 (CD, lower collecting duct), whereas 11ßHSD (CD, collecting duct) nearly completely co-localizes with mucin 1 . No spatial overlap can be detected for mucin 1 and megalin (PT, proximal tubule).Supplementary Figure 3: Localization of MUC1-fs in kidney tubules. Immunohistochemical analyses of serial human kidney sections for MUC1-fs and distinct kidney tubule markers (megalin, THP (Tamm Horsfall Protein), 11ßHSD (11-ß-hydroxysteroid dehydrogenase) and aquaporin-2. Partial co-localization is visible for MUC1-fs and THP (TAL, thick ascending limb) as well as for aquaporin-2 (CD, lower collecting duct), wheras 11 BHSD (CD, collecting duct) nearly completely colocalizes with MUC1-fs. No spatial overlap can be detected for MUC1-fs and megalin (PT, proximal tubule).

Supplementary Figure 4: Immunohistochemical validation of MUC1-fs antibodies. The four individual and purified antibodies against MUC1-fs (pAbx-fs) were used in immunohistochemistry on consecutive sections of one patient with ADTKD-MUC1 (ADTKD-0048). All antibodies produced identical positive signals, with pAb3-fs showing the most intense staining.

## SUPPLEMENTARY REFERENCES

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Supplementary Figure 1: Partial co-localization of mucin 1 and THP in kidney tubules


Supplementary Figure 2: Localization of mucin 1 in kidney tubule system


Supplementary Figure 3: Localization of MUC1-fs in kidney tubules


TAL
$a$ asi $\mathrm{a}^{\circ}$ \%

40x


Supplementary Figure 4: Immunohistochemical validation of MUC1-fs antibodies


