

Supplemental Figure 2

>lambda 1
QSALTQPMSGPGQSITISCTGT**NNDLGTYNF**VS^{WYQQHPGKAPKLVIYDVS}**YRPSGISYR**FSGSKSAYTASLTISDLQAGDEADYYC**TSFTTDSTFV**FGG
GTKLTVVGQPKAAPS^{VTLFPPSSEELQANKATLVCLISDFYPGAVTV}AWKADSSPVKAGVETTPSKQSNNKYAASSYLSLTPEQWKSHRSYS^{CQVTHEGSTV}
EKT
>lambda 2
SYVL^{TQAPS}S^VA^PG^QT^AR^IT^CE^GN**NIGSKS**AHWYQQRPGQAPVLVLY**DDSDRPSGIPER**FSGNSGN^TATL^ISGVEAGDEADFYC**QVWDNSNDHPV**FG
GG^{TTL}T^VL^GQ^PK^AAPS^VT^LF^{PPS}SEELQ^{AN}K^AT^LV^CLISDF^YPGAV^TAWKADSSPVKAGVETTPSKQSNNKYAASSYLSLTPEQWKSHRSYS^{CQVTHEGST}
VEKT
>lambda 3
YVL^{TQPS}S^VA^PG^QT^AR^IT^CGGH**NIGSKS**VHWYQQKPGQAPLLVVY**NNDRPSWIPER**FFGSNSGN^TATL^TVGVEAGDEADYYC**QIW**DSSGNPRV**VF**
GGGTKLTVLSQPKAAPS^VT^LF^{PPS}SEELQ^{AN}K^AT^LV^CLISDF^YPGAV^TAWKADSSPVKAGVETTPSKQSNNKYAASSYLSLTPEQWKSHRSYS^{CQVTHEGST}
TVEKT
>lambda 4
YVL^{TQPS}S^VA^PG^QT^AR^IT^CGGN**NIGYKS**VWHHQKPGQAPVLVY**DDSDRPSGIPER**FSGNSGN^TATL^TIRVEAGDEADFYC**QVWDSSSDQWV**FG
GGTKLTVLGQPKAAPS^VT^LF^{PPS}SEELQ^{AN}K^AT^LV^CLISDF^YPGAV^TAWKADSSPVKAGVETTPSKQSNNKYAASSYLSLTPEQWKSHRSYS^{CQVTHEGST}
VEKT
>kappa 1
IQMTQSPSSL^SASVGDRV^TITCRAS**SQSI**STS**LNWYQQ**KL^GK^AP^LLIY**AA**STL**QSGVPSRF**SGSASGTDFTLTISSLQ^{PED}FATYFC**QQSYSNSM**YTFGQG^{TK}
EIKRTVAAPS^VF^IF^{PPS}D^EQL^KSGTAS^VV^CLLNNF^YPREAKVQW^KVDNALQSGNSQESVTEQDSKD^TY^SLS^STL^TLSKADYE^EHKVYACEVTHQGLSSP^VT^K
FNR
>kappa 2
DIQMTQSPSSL^SASVGDRV^TITCQAS**QGIT**YLSWYQQRPGK^PK^LLIY**DA**V^NLES^GV^SSR^FSGSG^GT^DFTLTISSLQ^{PED}IGTYYC**QQYDSL**PPD^FGG^{TK}
VDIKRTVAAPS^VF^IF^{PPS}D^EQL^KSGTAS^VV^CLLNNF^YPREAKVQW^KVDNALQSGNSQESVTEQDSKD^TY^SLS^STL^TLSKADYE^EHKVYACEVTHQGLSSP^VT^K
SFNR
>kappa 3
DIQMTQSPSSL^SASVGDRV^TITCRAS**QGIS**NYLAFQQKPGK^AP^K**SLVY****A**SS^LQTG^VPSK^FSGSG^GT^DFTLTISSLQ^{PED}FATY^{YYC}**QQYNS**YP^TFGQ^{GT}
LEITRTVAAPS^VF^IF^{PPS}D^EQL^KSGTAS^VV^CLLNNF^YPREAKVQW^KVDNALQSGNSQESVTEQDSKD^TY^SLS^STL^TLSKADYE^EHKVYACEVTHQGLSSP^VT^K
SFNR
>kappa 4
IQMTQSPSSL^SASVGDRV^TITCQAS**EDIN**NY**LNWYQQ**KPGK^PK^VLIY**DA**S**N**LA^T**GVPL**RF^SSG^GGT^NFNFT^ISLQ^{PED}VAT^{YYC}**QQYNN**F^{PL}T^FGG^{GT}
RAEVKRTVAAPS^VF^IF^{PPS}D^EQL^KSGTAS^VV^CLLNNF^YPREAKVQW^KVDNALQSGNSQESVTEQDSKD^TY^SLS^STL^TLSKADYE^EHKVYACEVTHQGLSSP^VT^K
TKSFNR

Supplemental Figure 2. Eight Sanger sequenced light chains from eight MM patients. The red font indicates sequences corresponding to CDR1,2 and 3 as mapped by DomainGapAlign algorithm at the IMGT web portal. Tryptic peptides found by LC-MS/MS are indicated in bold font.