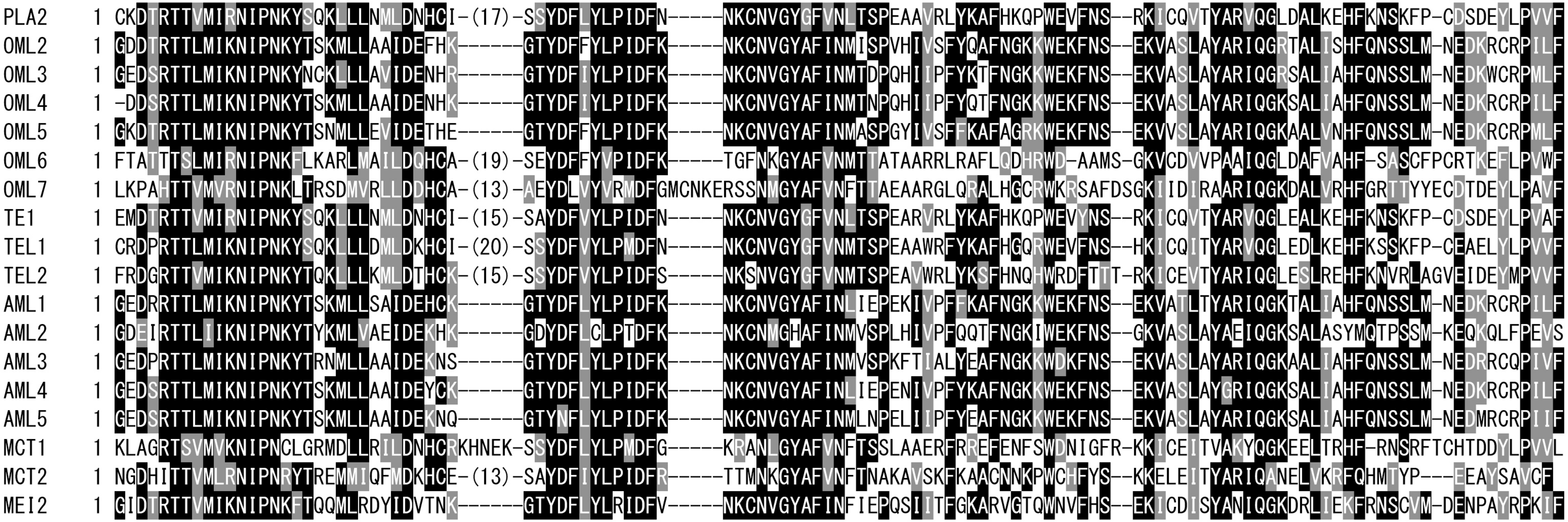

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TE1    1 EMDTRTTVMIRNIPNKYSQKLLLNMLDNHCI-(15)-SAYDFVYLPIDFN-----NKCNVGYGFVNLTSPEARVRLYKAFHKQPWEVYNS--RKICQVTYARVQGLEALKEHFKNSKFP-CDSDEYLPVAF
TEL1   1 CRDPRTTLMIKNIPNKYSQKLLLDMLDKHCI-(20)-SSYDFVYLPMDFN-----NKCNVGYGFVNMTSPEAAWRFYKAFHGQRWEVFNS--HKICQITYARVQGLEDLKEHFKSSKFP-CEAELYLPVVF
TEL2   1 FRDGRTTVMIKNIPNKYTQKLLLKMLDTHCK-(15)-SSYDFVYLPIDFS-----NKSNVGYGFVNMTSPEAVWRLYKSFHNQHWRDFTTT-RKICEVTYARIQGLESLREHFKNVRLAGVEIDEYMPVVF
AML1   1 GEDRRTTLMIKNIPNKYTSKMLLSAIDEHCK-------GTYDFLYLPIDFK-----NKCNVGYAFINLIEPEKIVPFFKAFNGKKWEKFNS--EKVATLTYARIQGKTALIAHFQNSSLM-NEDKRCRPILF
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AML3   1 GEDPRTTLMIKNIPNKYTRNMLLAAIDEKNS-------GTYDFLYLPIDFK-----NKCNVGYAFINMVSPKFTIALYEAFNGKKWDKFNS--EKVASLAYARIQGKAALIAHFQNSSLM-NEDRRCQPIVF
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MCT1   1 KLAGRTSVMVKNIPNCLGRMDLLRILDNHCRKHNEK-SSYDFLYLPMDFG-----KRANLGYAFVNFTSSLAAERFRREFENFSWDNIGFR-KKICEITVAKYQGKEELTRHF-RNSRFTCHTDDYLPVVL
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MEI2   1 GIDTRTTVMIKNIPNKFTQQMLRDYIDVTNK-------GTYDFLYLRIDFV-----NKCNVGYAFINFIEPQSIITFGKARVGTQWNVFHS--EKICDISYANIQGKDRLIEKFRNSCVM-DENPAYRPKIF
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Supplemental Figure 5