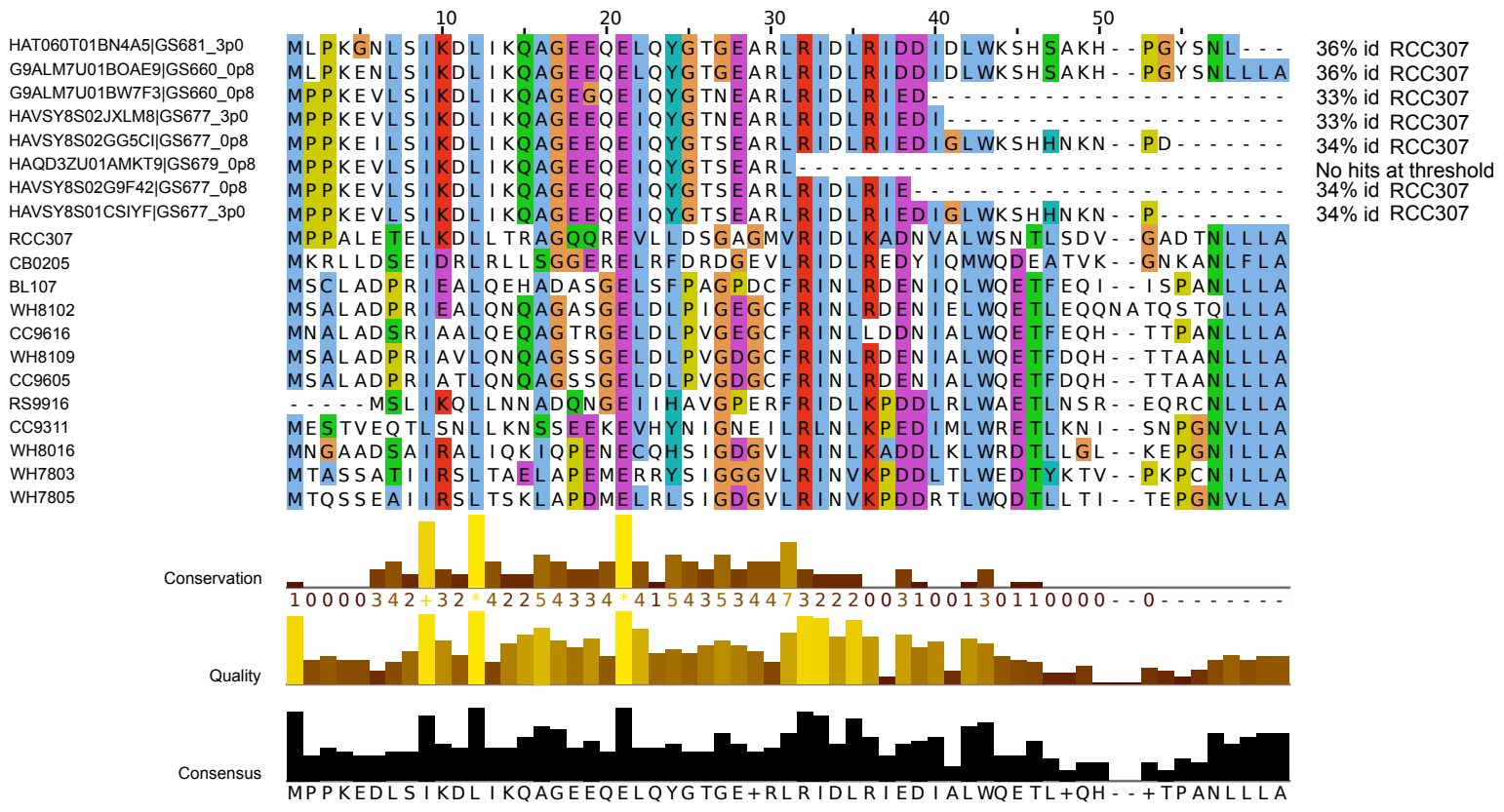
10
20
30
40
50
HAT060T01BN4A5|GS681_3p0 MLPKGNLSIKDLIKQAGEEQELQYGTGEARLRIDLRIDDIDLWKSHSAKH--PGYSNL--- 36% id RCC307
G9ALM7U01BOAE9|GS660_0p8 MLPKENLSIKDLIKQAGEEQELQYGTGEARLRIDLRIDDIDLWKSHSAKH--PGYSNLLLA 36% id RCC307
G9ALM7U01BW7F3|GS660_0p8 MPPKEVLSIKDLIKQAGEGQEIQYGTNEARLRIDLRIED---------- 33% id RCC307
HAVSY8S02JXLM8|GS677_3p0 MPPKEVLSIKDLIKQAGEEQEIQYGTNEARLRIDLRIEDI---------- 33% id RCC307
HAVSY8S02GG5CI|GS677_0p8 MPPKEILSIKDLIKQAGEEQEIQYGTSEARLRIDLRIEDIGLWKSHHNKN--PD------- 34% id RCC307
HAQD3ZU01AMKT9|GS679_0p8 MPPKEVLSIKDLIKQAGEEQEIQYGTSEARL---------- No hits at threshold
HAVSY8S02G9F42|GS677_0p8 MPPKEVLSIKDLIKQAGEEQEIQYGTSEARLRIDLRIE---------- 34% id RCC307
HAVSY8S01CSIYF|GS677_3p0 MPPKEVLSIKDLIKQAGEEQEIQYGTSEARLRIDLRIEDIGLWKSHHNKN--P------- 34% id RCC307
RCC307 MPPALETELKDLLTRAGQQREVLLDSGAGMVRIDLKADNVALWSNTLSDV--GADTNLLLA
CB0205 MKRLLDSEIDRLRLLSGGERELRFDRDGEVLRIDLREDYIQMWQDEATVK--GNKANLFLA
BL107 MSCLADPRIEALQEHADASGELSFPAGPDCFRINLRDENIQLWQETFEQI--ISPANLLLA
WH8102 MSALADPRIEALQNQAGASGELDLPIGEGCFRINLRDENIELWQETLEQQNATQSTQLLLA
CC9616 MNALADSRIAALQEQAGTRGELDLPVGEGCFRINLLDDNIALWQETFEQH--TTPANLLLA
WH8109 MSALADPRIAVLQNQAGSSGELDLPVGDGCFRINLRDENIALWQETFDQH--TTAANLLLA
CC9605 MSALADPRIATLQNQAGSSGELDLPVGDGCFRINLRDENIALWQETFDQH--TTAANLLLA
RS9916 -----MSLIKQLLNNADQNGEIIHAVGPERFRIDLKPDDLRLWAETLNSR--EQRCNLLLA
CC9311 MESTVEQTLSNLLKNSSEEKEVHYNIGNEILRLNLKPEDIMLWRETLKNI--SNPGNVLLA
WH8016 MNGAADSAIRALIQKIQPENECQHSIGDGVLRINLKADDLKLWRDTLLGL--KEPGNILLA
WH7803 MTASSATIIRSLTAELAPEMERRYSIGGGVLRINVKPDDLTLWEDTYKTV--PKPCNILLA
WH7805 MTQSSEAIIRSLTSKLAPDMELRLSIGDGVLRINVKPDDRTLWQDTLLTI--TEPGNVLLA
Conservation
10000342+32*42254334*41543534473222003100130110000--0--------
Quality
Consensus
MPPKEDLSIKDLIKQAGEEQELQYGTGE+RLRIDLRIEDIALWQETL+QH--+TPANLLLA