

	10	20	30	40	50	60
PRM2_RATTU	MVRYRMRSPS	ESP HQGPGQD	HESEEQG---	-QGQELNPER	VEDYGRTHRG	--HHRHRRCS
PRM2_MOUSE	MVRYRMRSPS	EGPHQGPGQD	HEREEQG---	-QGQGLSPER	VEDYGRTHRG	HHHHRHRRCS
PRM2_RATFU	MVRYRMRSPS	EGPHQGPGQD	HEREEQG---	-QGQELSPER	VEDYGRTHRG	--HHRHRRCS
PRM2_ALOSE	MVRYHVRSPS	ERPHEEYRQL	VNGQEQGRHG	QEEQGMSAEG	VEGYGRTHQG	CYGYRRRLCS
PRM2_CALJA	MVRYRVRSPS	ERPHEEYRQL	VNWQEQGRNG	QEEQGLSAEG	GEVYGRTHQG	YSSYRRRRCS
PRM2_SEMEN	MVRYRMRSLs	ERPHEVHGQQ	VYGQEQGHNG	QEEQGLSPEH	VEVYERTHQQ	YSHHRRRRCS
PRM2_ERYPA	MVRYRTRSLs	ERPHEVHGQQ	VHGQDQGHNG	QEEQGLSPEH	VEVYERTHQQ	HSHHRRRRCS
PRM2_MACNE	MVRYRMRSLs	ERPHEVHGQQ	VHGQDQGHNG	QEEQGLNPEH	VEVYERTHRG	HSHHRRRRCS
PRM2_MACFU	MVRYRMRSLs	ERPHEVHGQQ	VHGQDQGHNG	QEEQGLNPEH	VEVYERTHRG	HSHHRRRRCS
PRM2_MACMU	MVRYRMRSLs	ERPHEVHGQQ	VHGQDQGHNG	QEEQGLNPEH	VEVYERTH-G	HSHYRRRHCS
PRM2_GORGO	MVRCRVRSPS	ERSHEVYRQQ	LHGQEQGHHG	QEEQGLSPEH	VEVYERTH-G	HSHYRRRHCS
PRM2_PANPA	MVRYRVRSPS	EPSHEVYRQQ	LHGQEQGHHG	QEEQGLSPEH	VEVYERTH-G	HSHYRRRHCS
PRM2_PANTR	MVRYRVRSPS	EPSHEVYRQQ	LHGQEQGHHG	QEEQGLSPEH	VEVYERTH-G	HSHYRRRHCS
PRM2_HUMAN	MVRYRVRSLs	ERSHEVYRQQ	LHGQEQGHHG	QEEQGLSPEH	VEVYERTH-G	QSHYRRRHCS
PRM2_PONPY	MVRYCVRSLs	ERSHEVYRQQ	LHGQEQGHHD	QEEQGLSPEQ	VEVYERTQ-G	HSHYRRRHCS
PRM2_HYLLA	MVRYCVRSLs	ERSHEVYRQQ	LRGQEQGHHG	QEEQGLSPED	VEVYERTH-G	HSHYRRRHCS
	70	80	90	100	110	120
PRM2_RATTU	RKRLHRIHKR	R-RSCRRRRR	HSCCHRRRRH	RGCRRSRRRR	RCRCRKCRRH	CH
PRM2_MOUSE	RKRLHRIHKR	R-RSCRRRRR	HSCCHRRRRH	RGCRRSRRRR	RCRCRKCRRH	HH
PRM2_RATFU	RKRLHRIHKR	R-RSCRRRRR	HSCCHRRRRH	RGCRRSRRRR	RCKCRKCRRH	CH
PRM2_ALOSE	RRRLYRVHRR	RRSFCRRRC-	--CRYRRRR	RGCRTRRRRT	-----CRRH	--
PRM2_CALJA	RRRLYRVHRR	RRSFCRRRRR	RSCRYRRRPR	RGCRSRRRRR	-----CRRY	--
PRM2_SEMEN	RRRLYRVHRR	RHRSCRRRRR	RSCRHRRRRH	RGCRTRRRR	-----CRRY	--
PRM2_ERYPA	QRRLHRIHRR	RHRSCRRRRR	RSCRHRRRRH	RGCRTRRRR	-----CRRY	--
PRM2_MACNE	RRRLHRIHRR	RHRSCRRRRR	RSCRHRRRRH	RGCRTRRRR	-----CRRH	--
PRM2_MACFU	RRRLHRIHRR	RHRSCRRRRR	RSCRHRRRRH	RGCRTRRRR	-----CRRH	--
PRM2_MACMU	RRRLHRIHRR	RHRSCRRRRR	RSCRHRRRRH	RGCRTRRRR	-----CRRH	--
PRM2_GORGO	RRRLRRIHRQ	QHRSCRRRKR	RSCRHRRRRH	KGCRTRRRRT	-----CRRH	--
PRM2_PANPA	RRRLRRIHRQ	QHRSCRRRKR	RSCRHRRRRH	RGCRTRRRRT	-----CRKH	--
PRM2_PANTR	RRRLRRIHRQ	QHRSCRRRKR	RSCRHRRKHR	RGCRTRRRRT	-----CRRH	--
PRM2_HUMAN	RRRLHRIHRR	QHRSCRRRKR	RSCRHRRRRH	RGCRTRRRRT	-----CRRH	--
PRM2_PONPY	RRRLHRIHRQ	QHRSCRRRRR	HSCRHRRKHR	RGCRTRRRRT	-----CRRH	--
PRM2_HYLLA	RRRLHRIHRQ	QHRSCGRRRR	RSCRQRRRHR	RGCRTRRRR	-----CRRH	--

Supplemental Figure 3. Alignment of Eutherian P2 type sperm protamines.