		10 11 12 13 14 **15** 16 17 18 19 20 21 22
P33261	CP2CJ_HUMAN	CLSCL**L**LLSIWRQ
A0A2I3RMB5	A0A2I3RMB5_PANTR	CLSCL**L**LLSIWRQ
G3QCV6	G3QCV6_GORGO	CLSCL**L**LLSIWRQ
G1RRD4	G1RRD4_NOMLE	CLSCL**L**LLSLCRQ
A0A2R9B7Q9	A0A2R9B7Q9_PANPA	CLSCL**L**LLSIWRQ
		* * * * * * * * * * *

c.44T > A - p.L15H

Prediction pathogenic score

SIFT = 0 ; Damaging

MutPred = 0. 564 ; Pathogenic

PolyPhen-2 = 0.999 ; Probably Damaging

		399 400 401 402 403 404 **405** 406 407 408 409
P33261	CP2CJ_HUMAN	KEFPNP**E**MFDP
A0A2I3RMB5	A0A2I3RMB5_PANTR	KEFPNP**E**MFDP
G3QCV6	G3QCV6_GORGO	KEFPNP**E**MFDP
G1RRD4	G1RRD4_NOMLE	KEFPNP**E**MFDP
A0A2R9B7Q9	A0A2R9B7Q9_PANPA	KEFPNP**E**MFDP
		* * * * * * * * * * *

c.1215G > C - p.E405D

Prediction pathogenic score

SIFT = 0.169 ; Tolerated

MutPred = 0.425 ; Benign

PolyPhen-2 = 0.001 ; Bening