ERRATUM

Distribution and frequencies of *PDS* (*SLC26A4*) mutations in Pendred syndrome and nonsyndromic hearing loss associated with enlarged vestibular aqueduct: a unique spectrum of mutations in Japanese

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hum Pendrin	1	MAAPGGRSEF	10	20	30	40	50	60
mac Pendrin	1	MAAPGGRSEF	10	20	30	40	50	60
rat Pendrin	1	MAAPGGRSEF	10	20	30	40	50	60
hum Pendrin	51	ARSCSCSRKE	60	70	80	90	100	110
mac Pendrin	51	ARSCSCSRKE	60	70	80	90	100	110
rat Pendrin	51	ARSCSCSRKE	60	70	80	90	100	110
hum Pendrin	101	OGMAYALLAA	110	120	130	140	150	160
mac Pendrin	101	OGMAYALLAA	110	120	130	140	150	160
rat Pendrin	101	OGMAYALLAA	110	120	130	140	150	160
hum Pendrin	151	VVLSMAPDDE	160	170	180	190	200	210
mac Pendrin	151	VVLSMAPDDE	160	170	180	190	200	210
rat Pendrin	151	VVLSMAPDDE	160	170	180	190	200	210
hum Pendrin	201	LVPGGLQIGF	210	220	230	240	250	260
mac Pendrin	201	LVPGGLQIGF	210	220	230	240	250	260
rat Pendrin	201	LVPGGLQIGF	210	220	230	240	250	260
hum Pendrin	251	LSSTVTHET	260	270	280	290	300	310
mac Pendrin	251	LSSTVTHET	260	270	280	290	300	310
rat Pendrin	251	LSSTVTHET	260	270	280	290	300	310
hum Pendrin	301	PIRVIVTHIA	310	320	330	340	350	360
mac Pendrin	301	PIRVIVTHIA	310	320	330	340	350	360
rat Pendrin	301	PIRVIVTHIA	310	320	330	340	350	360
hum Pendrin	351	AASFSLAVVA	360	370	380	390	400	410
mac Pendrin	351	AASFSLAVVA	360	370	380	390	400	410
rat Pendrin	351	AASFSLAVVA	360	370	380	390	400	410
hum Pendrin	401	FVATTALSR	410	420	430	440	450	460
mac Pendrin	401	FVATTALSR	410	420	430	440	450	460
rat Pendrin	401	FVATTALSR	410	420	430	440	450	460
hum Pendrin	451	AAVIANLKE	460	470	480	490	500	510
mac Pendrin	451	AAVIANLKE	460	470	480	490	500	510
rat Pendrin	451	AAVIANLKE	460	470	480	490	500	510
hum Pendrin	501	GLPGGLQIGF	510	520	530	540	550	560
mac Pendrin	501	GLPGGLQIGF	510	520	530	540	550	560
rat Pendrin	501	GLPGGLQIGF	510	520	530	540	550	560
hum Pendrin	551	SSPIFYQNV	560	570	580	590	600	610
mac Pendrin	551	SSPIFYQNV	560	570	580	590	600	610
rat Pendrin	551	SSPIFYQNV	560	570	580	590	600	610
hum Pendrin	601	KNGIISLAV	610	620	630	640	650	660
mac Pendrin	601	KNGIISLAV	610	620	630	640	650	660
rat Pendrin	601	KNGIISLAV	610	620	630	640	650	660
hum Pendrin	651	PKVPIHSLV	660	670	680	690	700	710
mac Pendrin	651	PKVPIHSLV	660	670	680	690	700	710
rat Pendrin	651	PKVPIHSLV	660	670	680	690	700	710
hum Pendrin	701	ERLQCGQFT	710	720	730	740	750	760
mac Pendrin	701	ERLQCGQFT	710	720	730	740	750	760
rat Pendrin	701	ERLQCGQFT	710	720	730	740	750	760
hum Pendrin	751	DKCKPRLDE	760	770	780	790	800	810
mac Pendrin	751	DKCKPRLDE	760	770	780	790	800	810
rat Pendrin	751	DKCKPRLDE	760	770	780	790	800	810

Owing to a typesetting error, the sequence of Pendrin mutations was published incorrectly. The correct sequence is shown.

Furthermore, in Table 1 and Figure 2, IVS6-2 A–G mutation by Coucke *et al*¹⁷ should have been deleted.