

Table S1: Desmosomal disorders related to *JUP* mutation(s)

| Disease | Dermatological symptoms | | | | Cardiac involvement | Mode of inheritance | Mutation | Location | Total of patients | Total of families | Origin |
|--|-------------------------|-------------------|----------------|----------------------------|---------------------|---------------------|-----------------------|----------|-------------------|-------------------|--------------------------------|
| | PPK | Hair | Skin fragility | Others ectodermal features | | | | | | | |
| Naxos syndrome | + | woolly +/- sparse | - | - | ARVC | recessive | c.2157del2 | Exon 14 | 74 | NA | Greek / Turkish ¹⁻⁵ |
| CAPK syndrome | + | atrachia | - | - | ARVC | recessive | c.794G>A/ p.R265H | Exon 4 | 2 | 1 | Turkish ⁶ |
| Skin fragility + PPK + Woolly hair | + | woolly + sparse | + | - | no | recessive | c.71C>A/ p.S24X | Exon 2 | 3 | 1 | Argentina ⁷ |
| | | | | | | | c.468G>A | Exon 3 | 2 | 1 | Kuwaiti ⁷ |
| Lethal congenital epidermolysis bullosa | - | alopecia | + | onycholysis | no | recessive | c.1615C>T/ p.Q539X | Exon 9 | 1 | 1 | German ⁸ |

Sequences variations of *JUP* are referenced in the transcript GENE BANK accession NM_002230.

Table S2: Desmosomal disorders related to *DSP* mutation(s)

| Disease | Dermatological symptoms | | | | Cardiac involvement | Mode of inheritance | Mutation | Location | Total of patients | Total of families | Origin |
|--|-------------------------|----------------------|----------------|----------------------------|---------------------|---------------------|-------------------------------|----------|-------------------|-------------------|--------------------------|
| | PPK | Hair | Skin fragility | Others ectodermal features | | | | | | | |
| Striate PPK | + | - | - | - | no | dominant | c.939+1G>A | Intron 7 | 3 | 1 | NA ⁹ |
| | | | | | | | c.C991T/p.Q331X* | Exon 8 | 27 | 1 | NA ¹⁰ |
| Autosomal dominant Carvajal Syndrome | + | woolly and/or sparse | - | oligodontia | yes | dominant | c.1691C>T/p.T564I | Exon 13 | 3 | 2 | NA ^{11,12} |
| | | | | oligodontia | | | 30bp insertion | Exon 14 | 2 | 1 | NA ¹³ |
| | | | | oligodontia | | | c.1790C>T/p.S597L | Exon 14 | 3 | 1 | NA ¹⁴ |
| | | | | leuconychia | | | | | | | |
| | | | | nail dystrophy | | | c.1748T>C/p.L583P | Exon 14 | 1 | 1 | Caucasian ¹² |
| Autosomal recessive Carvajal Syndrome | + | woolly and/or sparse | - | - | yes | recessive | c.7901delG | Exon 24 | 12 | 3 | Ecuadorian ¹⁵ |
| | | | | | | | c.3799C>T/p.R1267X | Exon 23 | 1 | 1 | Turkish ¹⁶ |
| | | | | | | | c.6166G>C/p.G2056R | Exon 24 | 1 | 1 | Danish ¹⁷ |
| | | | | | | | c.3901C>T/p.Q1301X | Exon 23 | 1 | 1 | Indian ¹⁸ |
| | | | | | | | c.5208_5209delAG/p.G1737TfsX7 | Exon 24 | 2 | 1 | NA ¹⁹ |
| | | | | | | | c.7780delT/p.S2594PfsX8 | Exon 24 | 1 | 1 | Turkish ²⁰ |
| | | | | | | | c.7567delAAGA/p.K2523fsX37 | Exon 24 | 1 | 1 | NA ²¹ |
| c.6577G>A/p.E2193K | Exon 24 | | | | | | | | | | |

| | | | | | | | | | | | | |
|---|---|----------------------------|---|---|-----|-----------|--|---|--------------------|---|---|---------------------------|
| | | | | | | | | c.7566_7567delAAins C/ p.R2522SfsX39 | Exon 24 | 1 | 1 | NA ²² |
| | | | | | | | | c.7756C>T/ p.R2586X | Exon 24 | | | |
| | | | | | | | | c.1067C>A/ p.T356K | Exon 9 | 2 | 1 | NA ²² |
| | | | | | | | | c.2131_2132delAG/ p.S711CfsX4 | Exon 16 | | | |
| | | | | | | | | c.7123G>C/p.G2375R | Exon 24 | 9 | 1 | Israeli ²³ |
| | | | | | | | | c.3924delG/p. H1309TfsX1348 | Exon 23 | 1 | 1 | Palestinian ²⁴ |
| | | | | | | | | c.7111C>A, Q2371K | Exon 24 | 5 | 1 | Palestinian ²⁴ |
| Carvajal and skin fragility syndrome | + | woolly and/or sparse | + | - | yes | recessive | | c.4336C>T/p.Q1446X c.2017C>T/p.Q673X | Exon 23 Exon 15 | 1 | 1 | UK ²⁵ |
| | | | | | | | | c.4778_4790del13/p. L1593KfsX5 | Exon 23 | 1 | 1 | Swedish ²⁶ |
| | | | | | | | | c.6310delA/p.T2104 QfsX12 | Exon 24 | | | |
| | | | | | | | | c.4198C>T/R1400X c.6850C>T/R2284X | Exon 23 Exon 24 | 1 | 1 | Dominican ²⁷ |
| | | | | | | | | c.6310delA/ p.T2104QfsX12 | Exon 24 | 1 | 1 | Finnish ²⁸ |
| | | | | | | | | c.7964C>A/ p.A2655D | Exon 24 | | | |
| | | | | | | | | c.2516del4/p.H839fs X23 | Exon 18 | 1 | 1 | Brazilian ²⁹ |
| | | | | | | | | c.3971del4/p.N1324f | Exon 23 | | | |

| | | | | | | | sX23 | | | | |
|--|---------|----------------------|---|-----------------------------|---|-----------|------------------------|--------------------|---|---|----------------------|
| Skin fragility and woolly hair syndrome | + | woolly and/or sparse | + | nail dystrophy | no | recessive | c.2427T>A/p.C809X | Exon17 | 1 | 1 | NA ³⁰ |
| | | | | | | | c.861G>T/p.N287K | Exon 7 | | | |
| | | | | | | | c.1990C>T/p.Q664X | Exon 15 | 1 | 1 | NA ³⁰ |
| | | | | | | | c.7096C>T/p.R2366C | Exon 24 | | | |
| c.7097G>A/p.R2366H | Exon 24 | 5 | 1 | Saudi Arabian ³¹ | | | | | | | |
| c.7096C>T/p.R2366C | Exon 24 | 1 | 1 | Caucasian ³² | | | | | | | |
| c.6721_6722delAT/p.I2241FfsX3 | Exon 24 | | | | | | | | | | |
| Lethal acantholytic epidermolysis bullosa | | alope-cia | + | anonychia | poor cardiac contractility in one patient | recessive | c.5800C>T/p.R1934X** | Exon 24 Exon 24 | 1 | 1 | Dutch ³³ |
| | | | | | | | c.6370delTT | | | | |
| | | | | | | | c.2874del5/p.L959MfsX5 | Exon-Intron 20 | 2 | 1 | Yemeni ³⁴ |
| c.7248delT/p.F2416LfsX14 | Exon 24 | 1 | 1 | NA ³⁵ | | | | | | | |

Sequences variations of *DSP* are referenced in the transcript GENE BANK accession NM_004415.

c.C991T/p.Q331X* : this mutation was published as c.C1323T/p.Q331X. The correspondence between c.C1323T and p.Q331X was not found in ENSEMBL software. Therefore, we renamed it with this new combination.

c.5800C>T/p.R1934X** : This mutation was published as c.C6079T/p.R1934X . The correspondence between c.C6079T and p.R1934T was not found in ENSEMBL software. Therefore, we renamed it with this new combination.

Table S3: McGrath Syndrome, *PKP1* identified mutations

| Disease | Mode of inheritance | Mutation | Location | Total of patients | Origin |
|--|---------------------|--------------------------|-------------------------|-------------------|-------------------------|
| : Ectodermal Dysplasia/Skin fragility syndrome : - PPK - Skin fragility - Perioral lesions (at least 11 patients) - Sparse hair - Nail dystrophy - No cardiac involvement | recessive | c.203-1G>A | Intron 1 | 1 | Caucasian ³⁶ |
| | | c.213T>G/p.Y71X | Exon 2 | | |
| | | c.203-1G>A | Intron 1 | 3 | Arab ³⁷ |
| | | c.203-1G>T | Intron 1 | 2 | Egyptian ³⁸ |
| | | c.847-2A>G | Intron 4 | 1 | Arab ³⁷ |
| | | c.888delC | Exon 5 | 2 | Turkish ³⁹ |
| | | c.897del5/p.P299fsX61 | Exon 5 | 1 | Iraqi ⁴⁰ |
| | | c.910C>T/p.Q304X | Exon 5 | 1 | British ⁴¹ |
| | | c.1132ins28/PTC | Exon 6 | | |
| | | c.1054+1G>T | Intron 5 | 1 | Chinese ⁴² |
| | | c.1835-2A>G | Intron 10 | | |
| | | c.1233+2A>T | Intron 6 | 1 | French ⁴³ |
| | | c.1233-2A>G/p.R411SfsX51 | Intron 6 | 1 | Caucasian ⁴⁴ |
| IVS9+1G>A | Intron 9 | 1 | Dutch ⁴⁵ | | |
| c.2014C>T / p.R672X | Exon 11 | 3 | Brazilian ²⁹ | | |
| c.2021+1G>A | Intron 11 | 1 | Japanese ⁴⁶ | | |

Sequences variations of *PKP1* are referenced in the transcript GENE BANK accession NM_001005337.

Table S4: Desmosomal disorders related to *DSG1* and *DSG4* mutation(s)

| Disease | Gene | Mode of inheritance | Mutation | Location | Total of patients | Origin |
|---------------------|----------------------------|---------------------|-------------------|-----------|-------------------|-------------------------------------|
| Isolated PPK | <i>Desmoglein 1</i> | dominant | IVS2-1G>A | Intron 2 | 6 | Dutch ⁴⁷ |
| | | | c.76C>T/p.R26X | Exon 2 | 11 | Yemenite/pakistani ⁴⁸⁻⁵⁰ |
| | | | c.121insT/PTC | Exon 3 | 3 | Libyan ⁵¹ |
| | | | IVS4-2A>G | Intron 4 | 11 | Pakistani ⁵⁰ |
| | | | c.395C>A/p.S132X | Exon 5 | 3 | Iranian-Syrian ⁵² |
| | | | c.430A>T/p.R144X | Exon 5 | 4 | Scottish ⁵³ |
| | | | c.515C>T | Exon 5 | 10 | Pakistani ⁵⁰ |
| | | | c.C602T/p.Q201X | Exon 5 | 5 | Jewish Sephardic ⁵⁴ |
| | | | c.C655T/p.R219X | Exon 5 | 3 | Jewish Ashkenazi ⁵⁴ |
| | | | c.1079insC/PTC | Exon 9 | 1 | NA ⁴⁸ |
| | | | c.1095T>A/p.Y365X | Exon 9 | 1 | NA ⁴⁸ |
| | | | c.1189delA/PTC | Exon 9 | 1 | NA ⁴⁸ |
| | | | IVS9-3C>G | Intron 9 | 10 | Pakistani ⁵⁰ |
| | | | c.1399delA | Exon 10 | 3 | Pakistani ⁵⁰ |
| | | | c.1627delA/PTC | Exon 11 | 1 | NA ⁴⁸ |
| | | | c.1668-1G>T | Intron 11 | 11 | Pakistani ⁵⁵ |

| | | | | | | |
|---|---------------------|-----------|--|--------------------|----|------------------------------------|
| | | | c.1931delA /PTC | Exon 14 | 2 | Jewish Ashkenazi ⁵⁴ |
| SAM Syndrome : - Severe Skin dermatitis, - multiple Allergies - Metabolic wasting | <i>Desmoglein 1</i> | recessive | c.49-1G>A, in-frame skipping of exon 2 | Exon 2 | 2 | Arab ⁵⁶ |
| | | | c.1861delG/ p.A612Qfs.X3 | Exon 13 | 4 | Arab ⁵⁶ |
| | | | c.2659C>T/p.R887X | Exon 15 | 1 | NA ⁵⁷ |
| Localized Autosomal recessive Hypotrichosis (LAH Syndrome) | <i>Desmoglein 4</i> | recessive | c.Ex5_8del | Exon 5 to 8 | 34 | Pakistani ⁵⁸⁻⁶⁰ |
| | | | c.GG384-385TT/p.A129S | Exon 5 | 1 | Iraqi ⁶¹ |
| | | | c.87delG/PTC | Exon 3 | 6 | Pakistani ⁶² |
| Monilethrix | <i>Desmoglein 4</i> | recessive | c.574T>C/p.S192P c.2039insT/PTC | Exon 6 Exon 13 | 1 | Japanese ⁶³ |
| | | | c.799C>G/ P267R | Exon 7 | 9 | Iraqi ⁶⁴ |
| | | | p.P267R p.R289X | Exon 7 Exon 8 | 3 | Iraqi / Moroccan ⁶⁴ |
| | | | c.216+1G>T p.P267R | Intron 3 Exon 7 | 6 | Iraqi and Iranian ^{64,65} |
| | | | c.763delT c.216+1G>T | Exon 7 Intron 3 | 2 | Iranian ⁶⁴ |
| | | | p.P267R c.763delT | Exon 7 Exon 7 | 2 | Iraqi and Iranian ⁶⁴ |

| | | | |
|---|-------------------|---|------------------------|
| c.624delG/p.M208IfsX4 c.2468G>A/ p.W823X | Exon 6 Exon 16 | 1 | Japanese ⁶⁶ |
| c.2119delG/p.R707IfsX109 | Exon 14 | 1 | Japanese ⁶⁷ |

Sequences variations of *DSG1* and *DSG4* are referenced in the transcript GENE BANK accession NM_001942 and NM_177986 respectively.

Table S5: Desmosomal genodermatoses induced by *DSC2* and *DSC3* mutation(s)

| Disease | Gene | Mode of inheritance | Mutation | Location | Total of patients | Origin |
|--|----------------------|---------------------|-----------------------------|----------|-------------------|----------------------|
| - Woolly hair - Mild PPK - Left cardiac involvement | <i>Desmocollin 2</i> | recessive | c.1841delG/ p.S614fsX625 | Exon 12 | 2 | NA ⁶⁸ |
| Hypotrichosis | <i>Desmocollin 3</i> | recessive | c.2129T>G/p.L710X | Exon 14 | 4 | Afghan ⁶⁹ |

Sequences variations of *DSC2* and *DSC3* are referenced in the transcript GENEBANK accession NM_024422 and NM_001941, respectively.

Table S6: Desmosomal disorders related to *CDSN* mutation(s)

| Disease | Mode of inheritance | Mutation | Location | Total of patients | Total of families | Origin |
|---|---------------------|--|----------|---|-------------------|-----------------------------------|
| Hypotrichosis Simplex of the Scalp (HSS) | dominant | p.Q215X | Exon 2 | 26 (Israeli family) + 17 (spanish family) | 2 | Spanish and Israeli ⁷⁰ |
| | | p.Q200X | Exon 2 | US | 1 | Danish ⁷⁰ |
| | | p.Y239X* | Exon 2 | 42 | 1 | Mexican ⁷¹ |
| | | c.717C>G/p.Y239X* | Exon 2 | 3 | 1 | Chinese ⁷² |
| Peeling Skin Syndrome type B (PSS) | recessive | c.164_167dup GCCT/ p.T57PfsX6 | Exon 2 | 1 | 1 | Ashkenazi Jewish ⁷³ |
| | | c.175A>T/p.K59X | Exon 2 | 4 | 1 | German ⁷⁴ |
| | | c.424G>T/p.G142X | Exon 2 | 2 | 1 | NA ⁷⁵ |
| | | c.746delG/ p.G249VfsX40 | Exon 2 | 2 | 1 | NA ⁷⁶ |
| | | A genomic deletion at the PSORS1 locus removing the entire <i>CDSN</i> gene | | 1 | 1 | Japanese ⁷⁷ |
| | | A large homozygous deletion of 59,184bp extending from 40.6 kb upstream to 13.2 kb downstream of <i>CDSN</i> | | 1 | 1 | Japanese ⁷⁸ |

Sequences variations of *CDSN* are referenced in the transcript GENE BANK accession NM_001264.

* These two mutations seem to be the similar.

Legend of the 6 supplemental tables :

- = absent, + = present, NA = Not Available, ARVC = Arrhythmogenic Right Ventricular Cardiomyopathy, *CDSN* = Corneodesmosin gene, *DSC2* = Desmocollin 2 gene, *DSC3* = Desmocollin 3 gene, *DSG1* = Desmoglein 1 gene, *DSG4* = Desmoglein 4 gene, *DSP* = Desmoplakin gene, *JUP* = Plakoglobin gene, *PKP1* = Plakophilin 1 gene, PPK = Palmoplantar Keratoderma, PTC = Premature Termination Codon

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