

Diagnostic des maladies autoinflammatoires (Panel de 300 gènes)

ACP5	ADA	ADA2	ADAM17	ADAR	ADGRE2	AICDA	AIM2	AIRE	ALPI	ALPK1	ALPL	AP1S3	AP3B1	ARHGEF6
ARPC1B	ATAD3A	B2M	BACH2	BTB	C1QA	C1QB	C1QC	C1R	C1S	C2	C3	CARD11	CARD14	CARD9
CARMIL2	CASP1	CASP10	CASP8	CD27	CD3E	CD3G	CD40	CD40LG	CD70	CDC42	CEBPE	CFH	COG6	COL1A1
COL7A1	COPA	CORO1A	CTLA4	CYBA	CYBB	CYBC1	DCLRE1C	DDX58	DEF6	DKC1	DNASE1	DNASE1L3	DNASE2	DOCK2
DOCK8	DPP8	DPP9	EGFR	ELANE	ELF4	ELP1	EPCAM	ERBIN	ERCC4	ERCC6L2	F12	FAAP24	FAS	FASLG
FBLIM1	FCN3	FDP5	FERMT1	FOXP3	G3BP1	G6PC3	GALNT3	GATA2	GBP5	GSDMD	GUCY2C	HAVCR2	HPS1	HPS4
HPS6	IARS1	ICOS	IFIH1	IKBIP	IKBKB	IKBKG	IKZF1	IL10	IL10RA	IL10RB	IL16	IL18	IL1RAPL1	IL1RN
IL21	IL21R	IL27RA	IL2RA	IL2RB	IL2RG	IL36RN	IL6R	IL7R	IRAK1	IRAK4	ISG15	ITCH	ITGB2	ITK
JAK1	JAK2	KPNA1	KPNA2	KRAS	LACC1	LCK	LIG4	LIPA	LPIN2	LRBA	LSM11	LYN	LYST	MALT1
MAN2B1	MASP2	MCM3AP	MCM9	MEFV	MLKL	MPO	MRTFA	MSN	MVD	MVK	MYD88	NAE1	NCF1	NCF2
NCF4	NCKAP1L	NCSTN	NFAT5	NFIL3	NFKB1	NFKB2	NFKBIA	NGLY1	NLRC3	NLRC4	NLRP1	NLRP12	NLRP13	NLRP3
NLRP4	NLRP7	NOD1	NOD2	NOX1	NRAS	NSMCE3	OAS1	ORAI1	OTUD7B	OTULIN	PDGFRB	PGM3	PIK3CD	PIK3R1
PKN1	PKN2	PLCG2	PMVK	PNPT1	POLA1	POLE	POMP	PRF1	PRKCD	PSEN1	PSENE1	PSMA3	PSMB10	PSMB4
PSMB8	PSMB9	PSMG2	PSTPIP1	PSTPIP2	PTPN22	PYCARD	RAB27A	RAC2	RAG1	RAG2	RASGRP1	RASGRP3	RBCK1	RC3H1
RELA	RELB	RHOG	RIPK1	RIPK3	RNASEH2A	RNASEH2B	RNASEH2C	RND1	RNF31	RTEL1	SAA1	SAE1	SAMD9	SAMD9L
SAMHD1	SERPING1	SH2D1A	SH3BP2	SHARPIN	SKIV2L	SLC11A1	SLC27A1	SLC27A6	SLC29A3	SLC37A4	SLC7A7	SLC9A3	SLCO2A1	SOCS1
SPINK5	SPINT2	STAT1	STAT2	STAT3	STAT4	STAT5B	STIM1	STING1	STK4	STX11	STXBP2	STXBP3	SYK	TAP1
TAP2	TAPBP	TBK1	TERT	TET2	TGFB1	TGFB1	TGFB2	TNFAIP3	TNFRSF11A	TNFRSF13B	TNFRSF13C	TNFRSF1A	TNFRSF9	TNFRSF15
TRAF3IP2	TRAP1	TREM2	TREX1	TRIM21	TRIM22	TRNT1	TTC37	TTC7A	UBA1	UBA2	UBA3	UBA5	UBA6	UBA7
UBAC1	UBAC2	UNC13D	USP18	USP43	WAS	WASF2	WDR1	WIPF1	WNT2B	WNT6	XIAP	ZAP70	ZBTB24	ZNF341

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
ACP5	171640	259358	NM_001111035.2		Spondyloenchondrodysplasia with immune dysregulation (SPENCDI)	Récessive
ADA	608958	117771	NM_000022.3		Adenosine deaminase deficiency	Récessive
ADA2	607575	396032	NM_001282225.1		Deficiency in Adenosine deaminase 2 (DADA2)	Récessive
ADAM17	603639	299456	NM_003183.6		Neonatal inflammatory skin and bowel disease	Récessive
ADAR	146920	119502	NM_001111.5		Aicardi-Goutieres syndrome, Dyschromatosis symmetrica hereditaria	Récessive/ Dominante
ADGRE2	606100		NM_013447.3		Vibratory urticaria	Dominante
AICDA	605257	119554	NM_020661.3		Immunodeficiency with hyper-IgM type 2	Récessive
AIM2	604578		NM_004833.2			Candidat
AIRE	607358	119562	NM_000383.3		Autoimmune polyendocrinopathy syndrome , type I, with or without reversible metaphyseal dysplasia	Récessive/ Dominante
ALPI	171740		NM_001631.4		Inflammatory bowel disease, Behçet-like	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
ALPK1	607347		NM_025144.3		Retinal dystrophy, Optic nerve edema, Splenomegaly, Anhidrosis and migraine Headache syndrome (ROSAH)	Dominante
ALPL	171760	119640	NM_000478.5		Hypophosphatasia and Chronic recurrent multifocal osteomyelitis	Récessive/ Dominante
AP1S3	615781	402648	NM_001039569.1		AP1 deficiency-mediated pustular psoriasis	Dominante
AP3B1	603401	121374	NM_003664.4		Hermansky-Pudlak syndrome	Récessive
ARHGEF6	300267	168322	NM_004840.2			Candidat
ARPC1B	604223		NM_005720.4		Platelet abnormalities with eosinophilia and immune-mediated inflammatory disease	Récessive
ATAD3A	612316	469968	NM_001170535.2		Type I interferonopathy	<i>Dominante</i>
B2M	109700	317771	NM_004048.3		MHC class I deficiency with cutaneous granulomas	<i>Récessive</i>
BACH2	605394		NM_021813.3		BACH2-related immunodeficiency and autoimmunity (BRIDA)	Dominante
BTK	300300	119094	NM_000061.2		Agammaglobulinemia Isolates growth hormone deficiency type III	Liée à l'X
C1QA	120550	221162	NM_015991.3		C1q deficiency, Monogenic SLE	Récessive
C1QB	120570	221166	NM_000491.4		C1q deficiency, Monogenic SLE	Récessive
C1QC	120575	221168	NM_001114101.2		C1q deficiency, Monogenic SLE	Récessive
C1R	613785	360362	NM_001733.4		Ehlers-Danlos syndrome, Monogenic SLE	Récessive
C1S	120580	221171	NM_201442.3		C1s deficiency, Ehlers-Danlos syndrome, Monogenic SLE	Récessive
C2	613927	119106	NM_000063.5		C2 deficiency, Monogenic SLE	Récessive
C3	120700	160064	NM_000064.3		C3 deficiency, Monogenic SLE	Récessive
CARD11	607210	333084	NM_032415.5		Severe combined immunodeficiency with atopic dermatitis and increased IgE, B cell expansion	Récessive/ Dominante
CARD14	607211	304694	NM_024110.4		CARD14-mediated psoriasis	Dominante
CARD9	607212	225286	NM_052813.4		Familial Candidiasis, Lyn deficiency-associated autoimmune and inflammatory diseases	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>CARMIL2</i>	610859	470980	NM_001013838.2		CARMIL2-Immunodeficiency with veo-IBD	Récessive
<i>CASP1</i>	147678	470036	NM_001257118.2			Candidat
<i>CASP10</i>	601762	119179	NM_032977.3		Autoimmune lymphoproliferative syndrome type II (ALPS)	Dominante
<i>CASP8</i>	601763	138851	NM_001228.4		Autoimmune lymphoproliferative syndrome with immunodeficiency	Récessive
<i>CD27</i>	186711	356847	NM_001242.4		Lymphoproliferative syndrome	Récessive
<i>CD3E</i>	186830	159200	NM_000733.3		Immunodeficiency, SCID-like	Récessive
<i>CD3G</i>	186740	201284	NM_000073.2		Immunodeficiency	Récessive
<i>CD40</i>	109535	119244	NM_001250.5		Immunodeficiency with hyper-IgM	Récessive
<i>CD40LG</i>	300386	119249	NM_000074.2		Immunodeficiency with hyper-IgM	Liée à l'X
<i>CD70</i>	602840	470052	NM_001252.4		Lymphoproliferative syndrome	Récessive
<i>CDC42</i>	116952	470056	NM_001791.3		Neonatal onset of pancytopenia, autoinflammation, rash, and episodes of hemophagocytic lymphohistiocytosis (NOCARH) / Takenouchi-Kosaki syndrome	Dominante
<i>CEBPE</i>	600749	183890	NM_001805.3		Noncanonical autoinflammatory inflammasomopathy with defective neutrophil function (CAIN)	Récessive
<i>CFH</i>	134370	119363	NM_000186.3		Complement factor H deficiency, Familial HLH	Récessive
<i>COG6</i>	606977	365113	NM_020751.2		Congenital disorder of glycosylation (CGD) type IIi	Récessive
<i>COL1A1</i>	120150	120704	NM_000088.3		Infantile cortical hyperostosis	Dominante
<i>COL7A1</i>	120120	120738	NM_000094.3		Dystrophic epidermolysis bullosa	Récessive/ Dominante
<i>COPA</i>	601924	448689	NM_001098398.1		Autoimmune interstitial lung, joint, and kidney disease	Dominante
<i>CORO1A</i>	605000	179462	NM_001193333.2		Immunodeficiency	Récessive
<i>CTLA4</i>	123890	120878	NM_005214.5		CTLA-4 haploinsufficiency with autoimmune infiltration (CHAI)	Dominante
<i>CYBA</i>	608508	138719	NM_000101.3		Chronic granulomatous disease	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>CYBB</i>	300481	120935	NM_000397.3		Chronic granulomatous disease	Liée à l'X
<i>CYBC1</i>	618334		NM_001033046.3		Chronic granulomatous disease	Récessive
<i>DCLRE1C</i>	605988	121027	NM_001033855.2		Omenn Syndrome; Athabascan-type severe combined immunodeficiency	Récessive
<i>DDX58</i>	609631	422901	NM_014314.3		Singleton-Merten syndrome	Dominante
<i>DEF6</i>	610094		NM_022047.3		Immunodeficiency syndrome with systemic autoimmunity and aberrant CTLA-4 homeostasis	Récessive
<i>DKC1</i>	300126	121091	NM_001363.4		Dyskeratosis congenita	Liée à l'X
<i>DNASE1</i>	125505	353586	NM_005223.3		Monogenic systemic lupus erythematosus	Récessive/ Dominante
<i>DNASE1L3</i>	602244	291856	NM_004944.3		Monogenic systemic lupus erythematosus	Récessive/ Dominante
<i>DNASE2</i>	126350		NM_001375.2		Monogenic systemic lupus erythematosus	Récessive/ Dominante
<i>DOCK2</i>	603122	457458	NM_004946.3		Immunodeficiency	Récessive
<i>DOCK8</i>	611432	220922	NM_203447.3		Hyper-IgE recurrent infection syndrome (HIES)	Récessive
<i>DPP8</i>	606819		NM_130434.4			Candidat
<i>DPP9</i>	608258	353250	NM_139159.4		DPP9 deficiency (NLRP1-related inflammasomopathy)	Récessive
<i>EGFR</i>	131550	121311	NM_005228.4		Neonatal inflammatory skin and bowel disease 2	Récessive
<i>ELANE</i>	130130	121511	NM_001972.3		Cyclic neutropenia, severe congenital neutropenia	Dominante
<i>ELF4</i>	300775	306189	NM_001421.3		Deficiency in ELF4, X-linked (DEX)	Liée à l'X
<i>ELP1</i>	603722	122607	NM_003640.4		Dysautonomia, familial	Récessive
<i>EPCAM</i>	185535	226018	NM_002354.2		Congenital diarrhea with tufting enteropathy	Récessive
<i>ERBIN</i>	606944	470292	NM_001253697.1		ERBIN deficiency	Dominante
<i>ERCC4</i>	133520	121595	NM_005236.2			Candidat

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
ERCC6L2	615667	403281	NM_001010895.2		Bone marrow failure syndrome	Récessive
F12	610619	121663	NM_000505.3		Cold-induced Urticarial Autoinflammatory Syndrome Related to Factor XII Activation (FACAS)	Dominante
FAAP24	610884		NM_152266.4		EBV-driven lymphoproliferative disease	Récessive
FAS	134637	121733	NM_000043.5		Autoimmune lymphoproliferative syndrome (ALPS)	Récessive/ Dominante
FASLG	134638	121740	NM_000639.2		Autoimmune lymphoproliferative syndrome (ALPS)	Récessive/ Dominante
FBLIM1	607747		NM_017556.3		Monogenic CRMO	Récessive
FCN3	604973	332191	NM_003665.3		Immunodeficiency due to ficolin 3 deficiency	Récessive
FDPS	134629	444574	NM_002004.3		Porokeratosis	Dominante
FERMT1	607900	159172	NM_017671.4		Kindler syndrome	Récessive
FOXP3	300292	121913	NM_014009.3		Immunodysregulation, polyendocrinopathy and enteropathy (IPEX)	Liée à l'X
G3BP1	608431		NM_005754.2			Candidat
G6PC3	611045	171034	NM_138387.3		Severe Congenital neutropenia	Récessive
GALNT3	601756	122006	NM_004482.3		Hyperphosphatemic familial tumoral calcinosis with CRMO	Récessive
GATA2	137295	274222	NM_001145661.1		Immunodeficiency	Dominante
GBP5	611467		NM_052942.3			Candidat
GSDMD	617042		NM_024736.6			Candidat
GUCY2C	601330	317726	NM_004963.3		Familial Diarrhea	Dominante
HAVCR2	606652		NM_032782.4		T-cell lymphoma subcutaneous panniculitis-like	Récessive
HPS1	604982	122480	NM_000195.4		Hermansky-Pudlak syndrome	Récessive
HPS4	606682	122486	NM_022081.5		Hermansky-Pudlak syndrome	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>HPS6</i>	607522	122492	NM_024747.5		Hermansky-Pudlak syndrome	Récessive
<i>IARS1</i>	600709		NM_013417.3		Very early-onset IBD	Récessive
<i>ICOS</i>	604558	122565	NM_012092.3		Common variable immunodeficiency (CVID)	Récessive
<i>IFIH1</i>	606951	402570	NM_022168.4		Aicardi-Goutieres syndrome, Singleton-Merten syndrome	Dominante
<i>IKBIP</i>	609861		NM_153687.3			Candidat
<i>IKBKB</i>	300248	122614	NM_001556.3		Ectodermal dysplasia and immunodeficiency	Récessive/ Dominante
<i>IKBKG</i>	300248	122614	NM_003639.4		Ectodermal dysplasia and immunodeficiency NEMO deleted exon 5-autoinflammatory syndrome (NEMO-NDAS)	Liée à l'X
<i>IKZF1</i>	603023	318745	NM_006060.5		Common variable immunodeficiency (CVID), Monogenic SLE	Dominante
<i>IL10</i>	124092	122623	NM_000572.3		Early onset inflammatory bowel disease	Récessive
<i>IL10RA</i>	146933	244737	NM_001558.3		Early onset inflammatory bowel disease (IBD)	Récessive
<i>IL10RB</i>	123889	244742	NM_000628.4		Early onset inflammatory bowel disease (IBD)	Récessive
<i>IL16</i>	603035		NM_004513.5			Candidat
<i>IL18</i>	600953	470496	NM_001562.3			Candidat
<i>IL1RAPL1</i>	300206	159931	NM_014271.3			Candidat
<i>IL1RN</i>	147679	212902	NM_173842.2		Deficit of IL-1 receptor agonist (DIRA)	Récessive
<i>IL21</i>	605384	402379	NM_021803.4		IL-21 deficiency with early onset IBD	Récessive
<i>IL21R</i>	605383	357435	NM_021798.3		Immunodeficiency	Récessive
<i>IL27RA</i>	605350		NM_004843.3			Candidat
<i>IL2RA</i>	147730	160265	NM_000417.2		Immunodeficiency with lymphoproliferation and autoimmunity	Récessive
<i>IL2RB</i>	146710	355507	NM_000878.4		Immunodeficiency with lymphoproliferation and autoimmunity	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>IL2RG</i>	308380	122641	NM_000206.2		Severe Combined Immunodeficiency; Moderate Immunodeficiency	Liée à l'X
<i>IL36RN</i>	605507	281457	NM_173170.1		Deficit of IL-36 receptor agonist (DITRA)	Récessive
<i>IL6R</i>	147880		NM_000565.3		Hyper-IgE recurrent infection syndrome (HIES)	Récessive
<i>IL7R</i>	146661	159734	NM_002185.4		Severe combined immunodeficiency, T-cell negative, B-cell/natural killer cell-positive type	Récessive
<i>IRAK1</i>	300283	369336	NM_001569.3			Candidat
<i>IRAK4</i>	606883	122674	NM_001114182.2		Immunodeficiency	Récessive
<i>ISG15</i>	147571	315470	NM_005101.4		Mendelian susceptibility to mycobacterial disease with basal ganglia calcification (MSMD)	Récessive
<i>ITCH</i>	606409	232245	NM_031483.7		Autoimmune disease, multisystem with facial dysmorphism	Récessive
<i>ITGB2</i>	600065	122698	NM_000211.4		Leukocyte adhesion deficiency (LAD)	Récessive
<i>ITK</i>	186973	244711	NM_005546.3		Lymphoproliferative syndrome	Récessive
<i>JAK1</i>	147795		NM_002227.3		Autoinflammation, immune dysregulation and eosinophilia	Dominante
<i>JAK2</i>	147796	122727	NM_004972.3			Candidat
<i>KPNA1</i>	600686	470578	NM_002264.3			Candidat
<i>KPNA2</i>	600685		NM_002266.3			Candidat
<i>KRAS</i>	19007	122879	NM_033360.3		RAS-associated autoimmune leukoproliferative disease Lymphoproliferative RAS-MAPK signaling interferonopathy	<i>Somatique</i>
<i>LACC1</i>	613409	454001	NM_001128303.2		LACC1 deficiency, Juvenile arthritis	Récessive
<i>LCK</i>	153390	280151	NM_001042771.2		Immunodeficiency	Récessive
<i>LIG4</i>	601837	123058	NM_002312.3		LIG4 Syndrome	Récessive
<i>LIPA</i>	613497	123063	NM_001127605.2		Wolman disease (HLH)	Récessive
<i>LPIN2</i>	605519	123109	NM_014646.2		Majeed syndrome	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>LRBA</i>	606453	303168	NM_006726.4		Common variable immunodeficiency (CVID) with autoimmunity	Récessive
<i>LSM11</i>	617910		NM_173491.3		Aicardi-Goutieres syndrome	Récessive
<i>LYN</i>	165120		NM_002350.3		LYN associated autoinflammatory disease (LAID)	Candidat (Dominante)
<i>LYST</i>	606897	123318	NM_000081.3		Chediak-Higashi syndrome	Récessive
<i>MALT1</i>	604860	159756	NM_006785.4		Immunodeficiency	Récessive
<i>MAN2B1</i>	609458	123328	NM_000528.3		Metabolic IFNpathy	Candidat (Récessive)
<i>MASP2</i>	605102	332185	NM_006610.3		MASP2 deficiency	Récessive
<i>MCM3AP</i>	603294		NM_003906.4			Candidat
<i>MCM9</i>	610098	448630	NM_017696.2			Candidat
<i>MEFV</i>	608107	123191	NM_000243.2		Pyrin-associated autoinflammatory diseases (PAAD) including Familial Mediterranean fever (FMF) and Pyrin-associated autoinflammation with neutrophilic dermatosis (PAAND)	Récessive/ Dominante
<i>MLKL</i>	615153		NM_152649.3		PMID: 32561755	Candidat
<i>MPO</i>	606989	123468	NM_000250.1		Neutrophilic pustular skin disease	Candidat (Récessive)
<i>MRTFA</i>	606078	409876	NM_020831.4		Immunodeficiency	Récessive
<i>MSN</i>	309845	470694	NM_002444.2		Immunodeficiency	Liée à l'X
<i>MVD</i>	603236	444568	NM_002461.2		Porokeratosis	Récessive/ Dominante
<i>MVK</i>	251170	123588	NM_000431.3		Mevalonate kinase deficiency (MKD), Porokeratosis	Récessive/ Dominante
<i>MYD88</i>	602170	201114	NM_001172567.1		Immunodeficiency	Récessive
<i>NAE1</i>	603385		NM_003905.3		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
<i>NCF1</i>	608512	123695	NM_000265.5		Chronic granulomatous disease	Récessive
<i>NCF2</i>	608515	123701	NM_000433.3		Chronic granulomatous disease	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>NCF4</i>	601488	204415	NM_000631.4		Chronic granulomatous disease	Récessive
<i>NCKAP1L</i>	141180		NM_005337.4		Immunodeficiency with autoinflammation	Récessive
<i>NCSTN</i>	605254	268347	NM_015331.2		Hidradenitis suppurativa (HS)	Dominante
<i>NFAT5</i>	604708	470770	NM_138714.3		NFAT5 deficiency	Dominante
<i>NFIL3</i>	605327		NM_005384.2		NFIL3 deficiency	Récessive
<i>NFKB1</i>	164011	438054	NM_003998.3		Common variable immunodeficiency (CVID)	Dominante
<i>NFKB2</i>	164012	371170	NM_001077494.3		Common variable immunodeficiency (CVID)	Dominante
<i>NFKBIA</i>	164008	159788	NM_020529.2		Ectodermal dysplasia and immunodeficiency	Dominante
<i>NGLY1</i>	610661	406885	NM_018297.3		Congenital disorder of deglycosylation	Récessive
<i>NLRC3</i>	615648		NM_178844.3		PMID: 32447396	Candidat (Récessive)
<i>NLRC4</i>	605254	444736	NM_021209.4		Autoinflammation with infantile enterocolitis (AIEC), Familial cold autoinflammatory syndrome (FCAS)	Dominante
<i>NLRP1</i>	606636	139184	NM_033004.3		NLRP1- associated autoinflammation with arthritis and dyskeratosis (NAIAD)	Récessive/ Dominante
<i>NLRP12</i>	609648	138361	NM_144687.3		Familial cold autoinflammatory syndrome (FCAS)	Dominante
<i>NLRP13</i>	609660		NM_176810.2			Candidat
<i>NLRP3</i>	606636	123821	NM_001243133.1		NLRP3-associated autoinflammatory disease	Dominante
<i>NLRP4</i>	609648		NM_134444.4		PMID: 35467709	Candidat
<i>NLRP7</i>	609661	123834	NM_001127255.1		Ulcerative colitis and hydatidiform mole	Dominante
<i>NOD1</i>	605980		NM_006092.3			Candidat
<i>NOD2</i>	605956	123845	NM_022162.2		Blau syndrome	Dominante
<i>NOX1</i>	300225		NM_007052.4		NOX1 deficiency	Dominante

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>NRAS</i>	164790	221346	NM_002524.4		RAS-associated autoimmune leukoproliferative disease	Somatique
<i>NSMCE3</i>	608243		NM_138704.3		Lung disease, immunodeficiency, and chromosome breakage syndrome (LICS)	Récessive
<i>OAS1</i>	164350		NM_002534.3		OAS1-associated polymorphic autoinflammatory immunodeficiency (OPAID)	<i>Dominante</i>
<i>ORAI1</i>	610277	201295	NM_032790.3		Immunodeficiency due to ORAI1 deficiency	Récessive
<i>OTUD7B</i>	611748		NM_020205.3			Candidat
<i>OTULIN</i>	615712		NM_138348.5		OTULIN-related autoinflammatory syndrome (ORAS)	Récessive
<i>PDGFRB</i>	173410	138376	NM_002609.3		Penttinen syndrome	Dominante
<i>PGM3</i>	172100	446811	NM_001199917.1		PGM3-congenital disorders of glycosylation (CDG) with severe immunodeficiency	Récessive
<i>PIK3CD</i>	602839	400623	NM_005026.4		Immunodeficiency	Dominante
<i>PIK3R1</i>	171833	352217	NM_181523.2		Immunodeficiency	Récessive/ Dominante
<i>PKN1</i>	601032		NM_213560.1			Candidat
<i>PKN2</i>	602549		NM_006256.3			Candidat
<i>PLCG2</i>	600220	303819	NM_002661.4		Autoinflammation, antibody deficiency, and immune dysregulation syndrome (APLAID), Familial cold autoinflammatory syndrome (FCAS)	Dominante
<i>PMVK</i>	607622	444563	NM_006556.3		Porokeratosis	Dominante
<i>PNPT1</i>	610316	317560	NM_033109.4		Combined oxidative phosphorylation deficiency (Mitochondrial disease) / Deafness	Récessive
<i>POLA1</i>	312040	460832	NM_016937.3		X-linked pigmentary disorder, reticulate, with systemic manifestations	Liée à l'X
<i>POLE</i>	174762	356190	NM_006231.3		Facial dysmorphism-immunodeficiency-livedo-short stature syndrome (FILS)	Récessive
<i>POMP</i>	613386	281391	NM_015932.5		<i>POMP</i> -related autoinflammation and immune dysregulation disease (PRAID), <i>Proteasome-associated autoinflammatory syndrome (PRAAS)</i>	Dominante
<i>PRF1</i>	170280	117995	NM_005041.5		Familial hemophagocytic lymphohistiocytosis (HLH)	Récessive
<i>PRKCD</i>	176977	332077	NM_006254.3		Autoimmune lymphoproliferative syndrome (ALPS), type III / Monogenic SLE	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>PSEN1</i>	104311	118099	NM_000021.3		Hidradenitis suppurativa (HS)	Dominante
<i>PSENEN</i>	607632	268351	NM_172341.3		Hidradenitis suppurativa (HS)	Dominante
<i>PSMA3</i>	176843	470916	NM_002788.3		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSMB10</i>	176847		NM_002801.3		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSMB4</i>	602177	470918	NM_002796.2		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSMB8</i>	177046	258610	NM_148919.3		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSMB9</i>	177045	470920	NM_002800.4		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSMG2</i>	609702		NM_020232.4		Proteasome-associated autoinflammatory syndrome (PRAAS)	Récessive/ Dominante
<i>PSTPIP1</i>	606347	118114	NM_003978.4		<i>PSTPIP1</i> -associated autoinflammatory diseases (PAID)	Dominante
<i>PSTPIP2</i>	616046		NM_024430.3			Candidat
<i>PTPN22</i>	600716	317445	NM_015967.6			Candidat
<i>PYCARD</i>	606838		NM_013258.4			Candidat
<i>RAB27A</i>	603868	118188	NM_183235.2		Griscelli syndrome with HLH	Récessive
<i>RAC2</i>	602049	204411	NM_002872.4		Immunodeficiency with defective neutrophil chemotaxis	Récessive/ Dominante
<i>RAG1</i>	179615	118215	NM_000448.2		Combined cellular and humoral immune defects with granulomas	Récessive/ Dominante
<i>RAG2</i>	179616	118218	NM_000536.3		Combined cellular and humoral immune defects with granulomas	Récessive/ Dominante
<i>RASGRP1</i>	603962		NM_005739.3		Immunodeficiency	Récessive
<i>RASGRP3</i>	609531		NM_001139488.1			Candidat
<i>RBCK1</i>	610924	331658	NM_031229.3		Polyglucosan body myopathy with or without immunodeficiency (LUBAC deficiency)	Récessive
<i>RC3H1</i>	609424		NM_172071.4		Immune dysregulation and systemic hyperinflammation syndrome	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
<i>RELA</i>	164014	395954	NM_021975.3		Chronic mucocutaneous ulceration	Dominante
<i>RELB</i>	604758		NM_006509.3		Immunodeficiency	Récessive
<i>RHOG</i>	179505		NM_001665.3		RhoG deficiency	Récessive
<i>RIPK1</i>	603453		NM_003804.5		Cleavage resistant RIPK1-associated fever (CRAF), Immunodeficiency with autoinflammation	Récessive/ Dominante
<i>RIPK3</i>	605817		NM_006871.3			Candidat
<i>RNASEH2A</i>	606034	118329	NM_006397.2		Aicardi-Goutieres syndrome	Récessive
<i>RNASEH2B</i>	610326	118335	NM_024570.3		Aicardi-Goutieres syndrome	Récessive
<i>RNASEH2C</i>	610330	118340	NM_032193.3		Aicardi-Goutieres syndrome	Récessive
<i>RND1</i>	609038		NM_014470.3			Candidat
<i>RNF31</i>	612487	432144	NM_017999.4		HOIP deficiency (LUBAC deficiency)	Récessive
<i>RTEL1</i>	608833	330673	NM_016434.3		Pulmonary fibrosis and/or bone marrow failure	Dominante
<i>SAA1</i>	104750	330340	NM_001178006.2		Hereditary AA Amyloidosis	Dominante
<i>SAE1</i>	613294		NM_005500.2		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
<i>SAMD9</i>	610456	118460	NM_017654.3		Myelodysplasia, infection, growth restriction, adrenal hypoplasia, genital phenotypes, and enteropathy (MIRAGE) syndrome, Inherited bone marrow failure	Dominante
<i>SAMD9L</i>	611170	461215	NM_152703.4		SAMD9L-associated autoinflammatory disease (SAMD9L-SAAD)	Dominante
<i>SAMHD1</i>	606754	201570	NM_015474.3		Aicardi-Goutieres syndrome	Récessive
<i>SERPING1</i>	606860	118618	NM_000062.2		Hereditary Angioedema	Récessive/ Dominante
<i>SH2D1A</i>	300490	118679	NM_002351.4		Lymphoproliferative syndrome	Liée à l'X
<i>SH3BP2</i>	602104	118689	NM_003023.4		Cherubism	Dominante
<i>SHARPIN</i>	611885	471052	NM_030974.3		LUBAC deficiency	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
SKIV2L	600478	299641	NM_006929.4		Trichohepatoenteric syndrome	Récessive
SLC11A1	600266	158401	NM_000578.3			Candidat
SLC27A1	600691		NM_198580.2		Merkelsson-Rosenthal syndrome	Dominante
SLC27A6	604196		NM_014031.4		PMID: 32447396	Candidat (Récessive)
SLC29A3	612373	173057	NM_018344.5		Pigmentary hypertrichosis and non-autoimmune insulin-dependent diabetes (H syndrome)	Récessive
SLC37A4	602671	119474	NM_001164277.1		Glycogen storage disease (GSD)	Récessive
SLC7A7	603593	119709	NM_001126105.2		Lysinuric protein intolerance with HLH	Récessive
SLC9A3	182307	456093	NM_004174.3		Congenital diarrhea with secretory sodium	Récessive
SLCO2A1	601460	291830	NM_005630.2		Primary hypertrophic osteoarthropathy	Récessive
SOCS1	603597		NM_003745.1		Early-onset autoimmunity associated with SOCS1 haploinsufficiency / Monogenic systemic lupus erythematosus	Dominante
SPINK5	605010	119840	NM_001127698.1		Netherton syndrome	Récessive
SPINT2	605124	178801	NM_021102.3		Congenital diarrhea with secretory sodium	Récessive
STAT1	600555	119879	NM_007315.3		Immunodeficiency	Dominante
STAT2	600556	434238	NM_005419.3		Immunodeficiency with HLH	Récessive
STAT3	102582	138523	NM_139276.2		Autoimmune disease, multisystem, infantile-onset	Dominante
STAT4	600558	320815	NM_003151.3			Candidat
STAT5B	604260	119882	NM_012448.3		Hypereosinophilic syndrome with urticarial rash	Somatique
STIM1	605921	201300	NM_003156.3		Immunodeficiency	Récessive
STING1	612374	438066	NM_198282.3		STING-associated vasculopathy, infantile-onset (SAVI)	Dominante
STK4	604965	317798	NM_006282.4		T-cell immunodeficiency, recurrent infections, autoimmunity, and cardiac malformations (TIIAC)	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
STX11	605014	119892	NM_003764.3		Familial hemophagocytic lymphohistiocytosis (HLH)	Récessive
STXBP2	601717	227102	NM_006949.3		Familial hemophagocytic lymphohistiocytosis (HLH)	Récessive
STXBP3	608339		NM_007269.3		Very Early Onset Inflammatory Bowel Disease, Bilateral Sensorineural Hearing Loss and Immune Dysregulation	<i>Récessive/ Dominante</i>
SYK	600085		NM_003177.6		SYK-associated autoinflammatory disease	<i>Dominante</i>
TAP1	170260	123369	NM_000593.5		MHC class I deficiency with vasculitis or pyoderma gangrenosum	<i>Récessive</i>
TAP2	170261	119922	NM_000544.3		MHC class I deficiency with vasculitis or pyoderma gangrenosum	<i>Récessive</i>
TAPBP	601962	123371	NM_003190.4		MHC class I deficiency with vasculitis or pyoderma gangrenosum	<i>Récessive</i>
TBK1	604834	318865	NM_013254.3		TBK1 deficiency autoinflammation driven by TNF-induced cell death	Récessive
TERT	187270	138451	NM_198253.2		pulmonary fibrosis and Høyeraal-Hreidarsson syndrome	Dominante
TET2	612839	268325	NM_001127208.2		Immunodeficiency	Récessive
TGFB1	190180	120047	NM_000660.6		Camurati-Engelman disease	Dominante
TGFBR1	190181	120065	NM_004612.3		Loeys-Dietz syndrome, type 1	Dominante
TGFBR2	190182	120069	NM_001024847.2		Loeys-Dietz syndrome, type 2	Dominante
TNFAIP3	191163	369146	NM_006290.3		Haploinsufficiency in A20 or Familial autoinflammatory syndrome (HA20), Behcet-like	Dominante
TNFRSF11A	603499	120160	NM_003839.3		TRAPS-like disease	Dominante
TNFRSF13B	604907	120169	NM_012452.2		TACI deficiency	Récessive/ Dominante
TNFRSF13C	606269	231494	NM_052945.3		Immunodeficiency	Récessive
TNFRSF1A	191190	120173	NM_001065.3		TNF-receptor associated periodic fever syndrome (TRAPS)	Dominante
TNFRSF9	602250	471244	NM_001561.5			Candidat
TNFSF15	604052	317398	NM_005118.3			Candidat

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
TRAF3IP2	607043	370364	NM_147686.3		ACT1 deficiency, chronic mucocutaneous candidiasis	Récessive
TRAP1	606219		NM_016292.2		TRAP1 chaperone protein-associated autoinflammation	Récessive
TREM2	605086	120249	NM_018965.3		Familial hemophagocytic lymphohistiocytosis	Candidat
TREX1	606609	120255	NM_033629.5		Aicardi-Goutieres syndrome	Récessive/ Dominante
TRIM21	109092		NM_003141.3			Candidat
TRIM22	606559	471254	NM_006074.4		Granulomatous colitis and severe perianal disease	Récessive
TRNT1	612907	413615	NM_182916.2		Sideroblastic anemia with B-cell immunodeficiency, periodic fevers, and developmental delay (SIFD)	Récessive
TTC37	614589	238544	NM_014639.3		Trico hepato enteric syndrome	Récessive
TTC7A	609332	353460	NM_020458.3		Gastrointestinal defects and immunodeficiency syndrome	Récessive
UBA1	314370	140036	NM_003334.3		VEXAS syndrome	Somatique
UBA2	613295		NM_005499.2		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBA3	603172		NM_003968.3		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBA5	610552	469050	NM_024818.4		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBA6	611361		NM_018227.5		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBA7	191325		NM_003335.2		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBAC1	608129		NM_016172.2		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UBAC2		285499	NM_001144072.1		Groupe des Ubiquitin like modifier activating enzymes (UBA)	Candidat
UNC13D	608897	120416	NM_199242.2		Familial hemophagocytic lymphohistiocytosis (HLH)	Récessive
USP18	607057		NM_017414.3		Pseudo-TORCH syndrome	Récessive
USP43			NM_153210.4		USP43-associated autoinflammatory disease (USP43-AID)	Récessive

Gène	OMIM	Orphanet	Transcrit (RefSeq)	Code couleur Nouveaux gènes Gènes candidats de MAI	Phénotype	Hérédité
WAS	300392	120490	NM_000377.2		Wiskott-Aldrich syndrome	Liée à l'X
WASF2	605875		NM_006990.4			Candidat
WDR1	604734		NM_017491.4		Periodic fever, immunodeficiency, and thrombocytopenia syndrome (PFIT)	Récessive
WIPF1	602357	292448	NM_001077269.1		Wiskott-Aldrich syndrome	Récessive
WNT2B	601968		NM_024494.2		Diarrhea	Récessive
WNT6	604663		NM_006522.3			Candidat
XIAP	300079	138732	NM_001167.3		XIAP deficiency-associated lymphoproliferative syndrome	Liée à l'X
ZAP70	176947	120579	NM_001079.3		Immunodeficiency	Récessive
ZBTB24	614064	268103	NM_014797.2		Immunodeficiency-centromeric instability-facial anomalies syndrome	Récessive
ZNF341	618269		NM_032819.4		Hyper-IgE recurrent infection syndrome (HIES)	Récessive