Evidence for gliadin antibodies as causative agents in schizophrenia.

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Abstract

Antibodies to gliadin, a component of gluten, have frequently been reported in schizophrenia patients, and in some cases remission has been noted following the instigation of a gluten free diet. Gliadin is a highly immunogenic protein, and B cell epitopes along its entire immunogenic length are homologous to the products of numerous proteins relevant to schizophrenia (p = 0.012 to 3e-25). These include members of the DISC1 interactome, of glutamate, dopamine and neuregulin signalling networks, and of pathways involved in plasticity, dendritic growth or myelination. Antibodies to gliadin are likely to cross react with these key proteins, as has already been observed with synapsin 1 and calreticulin. Gliadin may thus be a causative agent in schizophrenia, under certain genetic and immunological conditions, producing its effects via antibody mediated knockdown of multiple proteins relevant to the disease process. Because of such homology, an autoimmune response may be sustained by the human antigens that resemble gliadin itself, a scenario supported by many reports of immune activation both in the brain and in lymphocytes in schizophrenia. Gluten free diets and removal of such antibodies may be of therapeutic benefit in certain cases of schizophrenia.

Introduction

A number of studies from China, Norway, and the USA have reported the presence of gliadin antibodies in schizophrenia ¹⁻⁵. Gliadin is a component of gluten, intolerance to which is implicated in coeliac disease ⁶. Both gluten intolerance and coeliac disease have also been associated with schizophrenia ^{7,8}, and remission of schizophrenia, in specific subsets of patients , has occasionally been reported following the instigation of a gluten-free diet ^{9, 10}. Gliadin is a polyglutamine repeat protein (Fig 1), and de facto, a homologue of the mutant polyglutamine proteins in Huntington's disease, Dentatorubropallidoluysian atrophy (DRPLA) , spinal and Bulbar Muscular Atrophy (Kennedy disease) and Spinocerebellar ataxias ¹¹. Gliadin antibodies have also been found in Huntington's disease and spinocerebellar ataxias and gluten per se has been associated with various forms of ataxia ^{12 13}. These studies may implicate gliadin in the pathology of these diseases. This is not the subject of this article.

As reported below, polyglutamine repeats are highly immunogenic, the more so with each addition of glutamine. A number of schizophrenia susceptibility gene products contain polyglutamine repeats, while others also display a high degree of homology to other regions of gliadin. Gliadin antibodies may thus play a role in the pathology of schizophrenia by cross-reactive targeting of key schizophrenia-related proteins.

Results

Gliadin is a polyglutamine (polyQ) repeat protein with an internal contiguous sequence of 22 glutamines (Fig 1). Gliadin is highly immunogenic and 181/296 (61%) of its residues are considered as B cell epitopes with the server-defined cut off index of 0.35 (Fig 2). http://www.polygenicpathways.co.uk/gliadin.htm. Polyglutamine repeats are also immunogenic, and all are above the threshold of 0.35 (Fig 2). The antigenicity increases with the number of glutamine repeats. The BLAST of gliadin (whole protein) versus the human proteome yielded 29 significant results, again including highly relevant proteins, many, but not all, influenced by the polyglutamine repeat (Table 1). These proteins belong to members of the DISC1 interactome and also include pre- and postsynaptic proteins related to glutamate, GABA, and neuregulin signalling. They are also involved in dopaminergic function, myelination, and dendritic spine development and to neurogenesis, inflammation and oxidative stress (Fig 3).

A more detailed analysis revealed an interesting type of homology that is common to many more proteins than are listed in Table 1. This is exemplified in Fig 4. The Clustal alignment of KCNN3 with gliadin shows a non-extensive homology with 18% amino acid identity shared by the two proteins, within the homologous region. The gliadin protein is characterised by many short repeat motifs, other than polyglutamines. These include (PQPQP) *4 ;(PQPQ) *5; (QPQP)*5; (PQQP)*2; (QQPY)*2; (QPQPQ)*2; (QQQQF)*2 and (VLQQ)*2. Some of these motifs can also be found in the KCNN3 protein, which, with the polyglutamine repeat contains 9 such areas of identity, many concentrated in a 13 amino acid contiguous tridecapeptide (QPQPPQLQQQQ). The overall gliadin/KCNN3 identity including these contiguous peptides is 5.3%. However, these matching gliadin/KCNN3 motifs extend over the whole length of the gliadin protein, which displays 31% identity with these KCNN3 fragments.

These fragments are mostly (12/15) within highly immunogenic regions of the gliadin protein. Because this immunogenicity extends over many different regions, several different antibodies to gliadin or to its partially digested fragments are likely to be produced. These repeat motifs and their presence in human proteins are likely to dramatically increase the likelihood of cross-reactivity between gliadin, gliadin fragments and human antigens.

These areas in human proteins were identified by sequential BLASTS of contiguous 25 amino acid fragments, along the length of the gliadin protein, each BLAST overlapping by 5 amino acids. By plotting these homologues along the length of the gliadin protein, these multiple repeats and their antigenicity could be identified simultaneously. In all, a total of 459 human proteins contain at least one of these tetrapeptide matches, 60 with pentapeptide matches, and others with longer contiguous matches as shown in Table 2.

This procedure allowed the definition of the number of gliadin matches within each human protein, and of the antigenic index of each of these matches. These results, for 5 or more matches are shown in Table 3: FOXP2, SMARCA2, NUMBL, KCNN3, and RAI1 contain from 14 to 22 of such matches. Certain key gene products including DISC1, neuregulin 2, dysbindin and synapsin 2 contain 5 or more of these gliadin consensus sequences. In fact, of the 459 proteins identified, 158 (34%) are the products of genes listed as susceptibility candidates in association studies. (See http://www.polygenicpathways.co.uk/gliadin.htm for details) An example of peptide matching to the antigenicity profile is shown in Fig 5 for KCNN3. This also allowed antigenicity mapping by family as shown in for glutamate related proteins or for myelin related proteins, many of which match gliadin in highly immunogenic regions (Fig 6).

A number of autoantibodies have been reported in schizophrenia. Their targets include the dopamine (DRD2) ¹⁴, glutamate (GRIN1) ¹⁵, acetylcholine (CHRNA7) ¹⁶ and opioid (OPRM1) receptors ¹⁷, heat shock protein 60 (HSPD1) ¹⁸ and hsp90 (HSP90A1) ¹⁹, MYC binding protein 2 (MYCBP2) ¹⁸ nerve growth factor (NGF) ²⁰ and striatin (STRN) ²¹ all of which are homologous to gliadin, with MYCBP2 and striatin particularly well represented (Fig 7). Myc inhibits myelination, and is also involved in dendrite and synapse formation ^{22, 23}. Striatin plays an important role in dendritic spine development ²⁴.

Discussion

The high immunogenicity of gliadin, over almost its entire length, suggests that multiple antibodies could be formed by presentation of diverse antigens in different cellular and tissue compartments. Such antibodies might be produced to the entire protein, or to its partially digested fragments. The homology with key schizophrenia related proteins, often covering many regions of gliadin is striking and suggests that gliadin antibodies could also target these human proteins. Indeed 9 of the autoantigens reported in schizophrenia patient are homologous to gliadin. It has also been shown that gliadin antibodies cross react with calreticulin and synapsin 1 ^{25, 26}. Synapsin 1 was homologous to the particular gliadin tested (PQQQP, PQQP and PLQQ) while calreticulin was not. It was however homologous to gamma gliadins from Triticum aestivum and Triticum urartu (VPRD), Triticum monococcum (VRPD) and a related goat grass, Aegilops searsii (PVIQ).

These homologous sequences are short, and in most cases, tetrapeptides, although higher degrees of homology were observed in many proteins (from 5 to 9 amino acids, not including

polyglutamine repeats (Table 2). Antibodies are quite capable of binding to such short epitopes ²⁷. In addition the repeat motifs in gliadin have already been noted and are, *per se*, immunogenic ²⁸.

The key pathological features of schizophrenia include reduced dendritic spine density and synaptic poverty ²⁹, deficits in myelination and oligodendrocyte cell loss ^{30, 31}, impaired neuregulin signalling ³²and imbalances in glutamate and dopamine neurotransmission ³³. Many gene products covering these networks are homologous to gliadin. The DISC1 network, connected to many of these areas³⁴ is also clearly targeted by gliadin homologues (Fig 1).

Antibodies are able to enter the brain via blood-brain barrier transporters ³⁵ and can also enter cells via a high affinity immunoglobulin receptor, tripartite motif-containing 21 (TRIM21) ³⁶.

This suggests that antibody related protein knockdown, of multiple proteins relevant to schizophrenia could be a direct consequence of gliadin allergy. Many studies have reported evidence of immune activation in schizophrenia patients, both in the brain ^{37, 38} and in lymphocytes ^{39, 40}, and autoimmune attack of certain cells may well explain some of the ongoing pathology of schizophrenia, for example oligodendrocyte ³⁰ and grey matter loss ⁴¹.

Following digestion, gliadin will be broken into peptide fragments that may also find their way into the brain via the circulation, and uptake via peptide transporters. As partial homologues of many relevant proteins, they may also be able to interfere with the signalling processes controlled by these proteins. Several pharmacological effects of gliadin or gluten have indeed been noted. For example gluten peptides have opioid activity ⁴²: Gliadin peptides are also able to activate protein kinase A ⁴³ and bind to the chemokine receptor CXCR3 ⁴⁴. They are also able to interfere with epidermal growth factor signalling in a number of cell lines ⁴⁵ and activate nuclear factor kappa beta signalling in monocytes ⁴⁶. This type of homology with human proteins may apply to allergens in general, whose deleterious effects are not necessarily restricted to immune activation, but also to the possibility of interaction with a multitude of host proteins that they resemble.

Similar protein matches are found in many proteins expressed by the viruses implicated in Alzheimer's disease ⁴⁷ or schizophrenia, and indeed these viral consensus sequences tend to be longer. For example hexapeptide identity between influenza viral proteins and several schizophrenia relevant proteins has been noted: These include reelin, neurexin 1-alpha and DISC1 ⁴⁸. In fact several hundred schizophrenia susceptibility gene products display this type of homology to diverse viruses and parasites (T.Gondii and B. Burgdorferri) implicated as risk factors in schizophrenia. It has recently been shown that DNA from many common non-retroviral viruses is integrated into mammalian genomes ⁴⁹. BLAST analyses of the human proteome also shows that thousands of human proteins contain these viral (or in this case allergen) contiguous matching sequences. Indeed this type of viral homology appears to cover the entire human genome. (See http://www.polygenicpathways.co.uk/blasts.htm). The human proteome has been estimated to contain ~33869 proteins with an average length of 375 amino acids ⁵⁰. For pentapeptide matches, this yields a figure of 370*33869 potential matching blocks (12.53 million). These building blocks are identical to those in viral, bacterial, fungal and allergen proteins. Upon

infection or ingestion, these pathogenic proteins are likely to seed havoc in the panoply of the host's signalling networks via the mechanisms described above, and are likely to contribute to the pathology of many human diseases.

It should also be noted that the viral and allergen protein homology is of course reflected at the DNA level and that viral double stranded DNA, plant or bacterial DNA is indistinguishable from our own. It is thus plausible that many gene association studies, using blood samples, have been indexing infection and ingestion as well as identifying key susceptibility genes. This is less of a problem when using long DNA probes and in no way detracts from the gene association results, whose relevance is generally supported by a plethora of experimental data related to the function of the genes identified. However, the confluence of gene and risk factor homology suggests that many genes are risk factors precisely because they encode for proteins that are homologous to those expressed by the viral, bacterial and allergen risk factors. However, they may only act as risk factors when such confluence is achieved, perhaps explaining the problems of replication in both gene and risk factor association studies.

Clearly gliadin, a major dietary component, cannot cause schizophrenia in all cases. Gluten intolerance and gliadin allergy are evidently related to the immune system. Many immune related susceptibility genes (few of which were encountered in this study) have been reported in association studies, including genome-wide association studies⁵¹. The high proportion of other types of schizophrenia susceptibility gene products related to gliadin suggests that those genes that encode for proteins with gliadin homology may be considered as risk factors if and when their products are homologous to a particular form of gliadin. Many different forms of gliadin, from diverse plant and bacterial species exist, as do many polymorphic genes, and their resultant differing protein sequences. The marriage of genes and risk factors, and the status of our immune system are thus three variables whose convergence may be obligatory to initiate the processes described above.

Gliadin antibodies and gluten intolerance have often been associated with schizophrenia, and in some cases a gluten free diet has been reported to evoke the remission of symptoms (see introduction). These data suggest that this is more than a simple association and that gliadin antibodies could well be the causative agent of schizophrenia in genetically and immunologically compromised individuals, and illustrate how this might be achieved. If so, then the instigation of a gluten free diet, as already shown ^{9, 10}, may be an effective substitute for drug related interventions. However in certain cases, even if gliadin is removed, the homologous human proteins might well be able to sustain the production of further antibodies, due to the permanence of autoantigens in the human biological network. Antigen and antibody removal by immunoadsorption techniques, or immunosuppression, might thus prove to be effective therapies. Ways of identifying the subsets of patients, who might benefit from such strategies, including routine antibody detection, may have a marked effect on the prevalence and severity of schizophrenia.

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Methods.

Gliadin is a polyglutamine (polyQ) repeat protein with an internal stretch of 22 glutamines. This sequence as well as gliadin or gliadin internal fragments were screened against the human proteome using the filters "schizophrenia", "glutamate", "dopamine" or "myelin" to trawl for proteins that might be related to gliadin (BlastP)⁵². Without these filters, many other polyglutamine proteins (Huntingtin, ataxins, the androgen receptor etc.) masked any underlying results. B-Cell epitopes within the gliadin protein were identified using the BepiPred server http://www.cbs.dtu.dk/services/BepiPred/, which predicts antigenicity related to the charge and hydrophobicity properties of the peptide ⁵³. The BLAST results and supplementary data can be visualised at http://www.polygenicpathways.co.uk/gliadin.htm where a NextBio highlighting tool provides details for all gene symbol abbreviations.

Table 1

Polyglutamine repeat proteins or proteins with significant overall homology to gliadin (BlastP gliadin vs. "schizophrenia", "glutamate", "dopamine" or "myelin"). Brief descriptions of function are provided for each protein. The number of glutamines is represented by, for example, Q22, or by the Q containing sequence in the human protein. E values are provided for the BLAST: If none is shown, the results were not significant (P> 0.05)

Proteins	QN and /or BLAST E value	
Glutamate and	transmission	
GRINA glutamate receptor, ionotropic, N- methyl D-aspartate- associated protein 1	E value 2e-11	Uncharacterise d glutamate binding NMDA receptor subunit
GRIN3B glutamate receptor, ionotropic, N- methyl-D- aspartate 3B	Q7	NMDA receptor subunit expressed in the human hippocampus and neocortex ⁵⁴
GRM1 glutamate receptor, metabotropic 1	Q5	Metabotropic glutamate receptor
IQSEC2 IQ motif and Sec7 domain 2	No polyglutamines BLAST E value 0.016	Part of the post-synaptic density, binding to psd95 (DLG4) ⁵⁵
NOS1AP : nitric oxide synthase 1 (neuronal) adaptor protein	HHQMQLLQQLLQQQQTQ	Plays a role in both presynaptic and postsynaptic (NMDA)

		glutamatergic function and controls dendritic spine development ^{56,} ⁵⁷
RIMS2 : regulating synaptic membrane exocytosis 2	QQFEMYKEQVKKMGEESQQQQEQ	Regulates calcium dependent neurotransmitte r release ⁵⁸
SYN3 : synapsin III	QQ RLSP Q G QQ PLSP Q SGSP QQQ	Presynaptic protein regulating neurotransmitte r release ⁵⁹
GRIPAP1 GRIP1 associated protein 1	QQQQEQEEALKQ-	Part of the postsynaptic scaffold for glutamate (AMPA) receptors ⁶⁰
MAP1A : microtubule- associated protein 1A	QQTHEQQQQ	Part of the NMDA receptor postsynaptic density, that controls dendrite branching and synapse formation ⁶¹ ; Binds to postsynaptic density proteins that connect to NMDA Kainate , AMPA and ERBB receptors
SPTAN1 : spectrin,	QQQQ	Part of the postsynaptic
alpha, non-		density that

erythrocytic 1		regulates
(alpha-fodrin)		glutamate
		receptor
		signalling 62
RORB : RAR-	QKHQQRLQEQRQQQ	Little is known
related orphan		about this
receptor B		protein:
		However it
		binds to MAP6,
		a microtubule
		protein that
		controls
		synaptic
		organisation, in
		particular of
		glutamatergic
		synapses where it
		controls the
		expression of
		the glutamate
		transporter and
		presynaptic
		genes,
		synaptophysin
		and GAP-43,
		spinophilin and
		MAP2 ⁶³ .
Gamma amino	butyric acid (GABA) related	
ATF5 cyclic	No polyglutamines	Regulates the
AMP-		development of
dependent	0.008	neurones,
transcription		astrocytes and
factor ATF-5		oligodendrocyt
		es ⁶⁴
DGCR6 :	QKHQEAQQACRPHNLPVLQAAQQ	Little is known
DiGeorge		except that is
syndrome		binds to the
critical region		GABA-b
gene 6	BLAST E value 0.001	receptor
30110 0		GABBR1 ⁶⁵
SP4 : Sp4	QNAQDQSNSLQQVQIVGQPILQQIQIQQPQQQ	Regulatesgluta
transcription		mate

	1	
factor	BLAST E = 9e-05	(GRIN2A), GABA (GABBA4)
		(GABRA4)
		receptors (and
		many others):
		Also controls
		dendritic function ⁶⁶⁻⁶⁸
		function ** **
Dopamine and	serotonin related	· · · ·
FOXP2	Q22	Controls
Forkhead box	DIACT 20.25	pathways
P2	BLAST 3e-25	involved in
		embryonic and
		nervous-
		system
		development,
		neurogenesis,
		cell migration
		and cell death.
		FOXP2
		knockout mice
		have lower
		cerebral
		dopamine
		concentrations.
		Expressed in
		dopamine and
		cyclic
		adenosine
		3',5'-
		monophosphat
		e-regulated
		phosphoprotein
		containing
		neurones in the
		cerebral cortex
		FOXP2 also
		controls
		dendritic spine
		development
		and synaptic
		plasticity. 69-72
LMX1A : LIM		LMX1A and B
homeobox		control the

transprintion		day alanmast of
transcription		development of
factor 1, alpha		midbrain
LMX1B : LIM	HQQQQE QQ	dopamine
homeobox		neurones 73
transcription		
factor 1, beta		
KCNN3	Q19	Regulates the
potassium		activity of
intermediate/s	BLAST E Value: 2e-22	dopamine and
mall		serotonin
conductance		neurones and
calcium-		plays a role in
activated		glutamate-
channel,		mediated long
subfamily N,		term
member 3		potentiation ^{74,}
		75
MAGEL2	QQAQASGPQ	MAGEL2 and
MAGE-like		necdin bind to
protein 2	BLAST E Value 2e-05	FEZ1 and are
		involved in the
		control of
		axonal growth
		and in the
		development of
		brain
		dopamine,
		serotonin and
		noradrenaline
		neurones ⁷⁶
NALCN :	QQQSCSIIHSLRESQQQE	Sodium
sodium leak		channel
channel, non-		activated by
selective		reducing
		extracellular
		calcium levels:
		It is activated
		by substance P
		and
		neorotensin in
		ventral
		tegmental and
		hippocampal

		neurones 77
PDCD6IP programmed cell death 6 interacting protein	E value 6e-06	Apoptosis inhibitor that also binds to D1 and D3 dopamine receptors ⁷⁸
POU3F2 POU class 3 homeobox 2 brain-specific homeobox/PO U domain protein 2	Q18 E value = 4e-15	Controls the expression of neural progenitors and of the tryptophan hydroxylase gene (TPH2) ^{79,} ⁸⁰
PPP1R3F : protein phosphatase 1, regulatory (inhibitor) subunit 3F	QQQLPQLEPQ	Protein phosphatase 1 regulates the activity of many transmitter systems (glutamate, GABA, dopamine . serotonin, inter alia): It also modifies dendritic spine development . 81-86
RANBP9 RAN binding protein 9	Q8	Binds to the D1 dopamine receptor as well as to nerve growth factor receptors NGFR and trkB (NTRK2), androgen and glucocorticoid receptors and DISC1 ⁸⁷⁻⁹¹

SMARCA2 : SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	Q2 BLAST E value 3e-15	Binds to the promoter of the noradrenaline transporter SLC6A2 : It also controls neuronal development on the mid hindbrain ^{92, 93}
SMARCC1 : SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	QQMEQQQHGQNPQQAHQH And EQQRQQLLTERQNFHMEQLKYAELRARQQMEQQQHG QNPQQ	Regulates the function of the glucocorticoid receptors (NR3C1), and Nurr77 (NR4A1) ⁹⁴ both of which play an important role in the control of dopamine neurones ⁹⁵
Neuregulin , No	OTCH and growth	
NRG2 : neuregulin 2	QQQQQQREEQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQ	This particular neuregulin is localised to dendrites and controls the expression of DISC1 ; It also controls free radical release from microglia ⁹⁶⁻⁹⁸
NUMBL numb homolog (Drosophila)- like	Q20 BLAST E value 6e-13	Important regulator of cortical neurogenesis ^{99, 100}
RAI1 retinoic acid induced 1	Q14	Regulates BDNF expression and

	BLAST E Value = 4e-10	also regulates autoimmunity by inhibiting lymphocyte activation and and triggen receptor signalling ^{101, 102}
Dendritic spine	es and plasticity	
HDAC4 :	QQQHQQFLEKHKQQFQQQQLQ	Histone
histone deacetylase 4	BLAST E value 0.006	deacetylases regulate gene transcription by
HDAC5 histone deacetylase 5	QQQEMLAAKQQQEMLAAKRQQELEQQRQREQQRQ EELEKQRLEQQ BLAST E value 5e-05	modifying histones. HDAC4 is localised in the brain and shuttles
HDAC9 : histone deacetylase 9	QQQQQIQKQLLIAEFQKQHENLTRQHQAQLQEHIKLQQ ELLAIKQQQ	between the nucleus and cytoplasm in response to neuronal activity. HDAC4 is also localised in dendritic spines associated with the postsynaptic density: HDAC9 also shuttles from nucleus to cytoplasm in response to neuronal activity, and controls dendritic spine density ¹⁰³⁻¹⁰⁵
MCF2 : MCF.2 cell	QQDQLTERDKFQISLQQNDEKQQ	Controls dendritic spine

line derived transforming sequence		development
MED25 mediator complex subunit 25	VQQQ BLAST E Value 0.001	Regulates the retinoid receptor RAR , which controls dendritic spine formation, ¹⁰⁷
NR4A3 : nuclear receptor subfamily 4, group A, member 3 also known as nor-1	QQQHQQ	Regulates pyramidal neurone survival and hippocampal axon guidance and controls neurite extension ^{108,} 109
PURA : purine-rich element binding protein A	EQLHQQQQQQEE BLAST E value 0.014	A single stranded DNA (e.g. viral) binding protein involved in the transport of RNA to dendrites. Plays and important role in postnatal brain development and also controls the transcription of myelin basic protein in oligodendrocyt es ¹¹⁰⁻¹¹²
Myelin related		
FOXJ3 forkhead box	QQQP	Involved in odendrocyte

J3	BLAST E value 5E-0.5	liferation 113
NCOR2 nuclear receptor corepressor 2	Q17 BLAST E value = 6E-11	Controls thyroid hormone, retinoic acid and steroid receptors ¹¹⁴ . Retinoids play a key role in dopamine function, dendritic spine development and oligodendrocyt e differentiation ¹¹⁵⁻¹¹⁷
PEG10 paternally expressed 10	No Polyglutamines BLAST E value 0.006	Controls the expression of myelin basic protein ¹¹⁸
SGCE : sarcoglycan, epsilon	QTQQNLPHQTQIPQQQ	Sarcoglycans are typically associated with muscle cells: However SGCE is expressed in Schwann cells, and in monoaminergic neurones and is believed to regulate dopaminergic activity ¹¹⁹⁻¹²¹
Development a	and neurogenesis	I
DISC1 : disrupted in schizophrenia	QQLRREIEEQEQQLQ	Key "hub gene" connected to many schizophrenia

1		networks ³⁴
MACF1 : microtubule- actin crosslinking factor 1	QLQQLQSQLAHQTEQKTLQKQQNTCHQQ	MACF1 knockdown impairs cortical and other brain area development : It also plays a role in growth cone advance 122, 123
PCM1 : pericentriolar material 1	QQQQRELKQLQEE	Involved in neuronal migration and neurogenesis
PCNT : pericentrin	QQQALHSQQQ	Regulates the neuronal progenitor cells in the developing cortex ¹²⁵
SEL1L3 : sel- 1 suppressor of lin-12-like 3	QQQQQ	Little is known, except that SEL1L3 binds to tubulin TUBB3 which is involved in cortical development and neuronal migration ¹²⁶
Endocytosis		
CHERP calcium homeostasis endoplasmic reticulum protein	Q12 BLAST E value 7e-08	No data: Endoplasmic reticulum location
CLINT1 : clathrin	QQQNMQQ	Involved in clathrin

interactor 1		mediated
		endocytosis
		and the
		endosomal
		transport of
		SNARE
		proteins
		including STX7
		127
STX7 :	QQLQQKQQ	Involved in
syntaxin 7		endosomal to
		lysosomal
		traffic and in
		phagocytosis.
		Also expressed
		in neutrophils
		and
		macrophages
		128-131
Apoptosis and	l oxidative stress	
EP400 E1A	Q22	Involved in the
binding		regulation of
protein p400	BLAST E value 1E-16	oxidative stress
protoni proto		132
HIPK3 :	QQKLTSAFQQQH	Involved in the
homeodomain		Fas apoptosis
interacting		pathway, and
protein kinase		also controls
3		the expression
3		of cytochrome
		p450 CYPA11,
		the enzyme
		responsible for
		pregnelonone
		synthesis, the
		firsts step in
		steroid
		hormone
		production ^{133,} 134
SETD2 : SET	QREAQKQQQQMQ	Involved in
domain		embryonic
containing 2		vascular

(a histone methyltransfer ase)		development
SHISA5 protein shisa- 5 precursor	No polyglutamines BLAST E value e-04	Localised in the endoplasmic reticulum and nuclear membrane and involved in p53 and p73 mediated apoptosis ¹³⁶⁻¹³⁸
TFAP4 transcription factor AP-4	E value 0.002	AP4 controls the expression of p21 a cell cycle cyclin dependent kinase inhibitor induced by p53 ¹³⁹
Immune syster	n	
HLX H2.0- like homeobox	Q8 BLAST E value 6e-06	Regulates interferon gamma production in natural killer cells ¹⁴⁰
HOXA10 homeobox A10	PPQQQPPPPPQP E value 0.012	Part of a vitamin D3 dependent pathway regulating haematopoesis
ITIH4 : inter- alpha (globulin) inhibitor H4 (plasma Kallikrein- sensitive	QQQ KSPE QQ E	Acute phase protein involved in inflammatory processes ¹⁴²

glycoprotein)		
NFAT5 nuclear factor of activated T- cells 5, tonicity- responsive	Q17 BLAST E value 1e-16	Plays an important role in the immune system and in water homoeostasis : Polydipsia (excessive water drinking) and water imbalance have been observed in schizophrenia 143, 144
Translation init	tiation	
ABCF1 ATP- binding cassette, sub- family F (GCN20), member 1	Q10 BLAST E value 9E-04	Regulates translation initiation ¹⁴⁵
EIF4G3 eukaryotic translation initiation factor 4 gamma, 3	No polyglutamines BLAST E value 0.008	Translation initiation factor
Miscellaneous		
ALX4 ALX homeobox 4	Q4 BLAST E value 9e-08	Involved in cranial bone development ^{66,} 67, 109, 146, 147
ARID3B AT rich interactive domain 3B (BRIGHT-like)	Q11 BLAST E value 1e-05	No data
PHC2 polyhomeotic homolog 2	VIQQQPQPQQQQPPPQQ BLAST E value 6E-0.6	No data

Table 2 Proteins with hexapeptide, or greater, matches to the gliadin protein. The consensus sequence is shown on the left followed by the gene symbols. N is the number of contiguous amino acids : DISC1 (N=5) is included for interest: Q = glutamine, followed by the number of Q's e.g. Q22. Full tables with abbreviations and data provided by a NextBio highlighting service are available at http://www.polygenicpathways.co.uk/gliadin.htm

Sequence and Gene symbol	Ν
Q22 FOXP2 SMARCA2	22
Q20 NUMBL	20
Q19 KCNN3	19
Q18 POU3F2	18
Q14 RAI1	14
TAVR-PVP PRODH (Proline dehydrogenase)	9
Q8 NRG2	8
LQQ-QQQ NOS1AP KCNN3	7
PSSSLQQ ZDHHC8 (Zinc finger protein)	7
Q7 PURA	7
QP+RQVP FXR1 (Fragile X protein)	7
FPPQ+P CABIN1 (calcineurin binding protein)	6
P+PQQP SYN2	6
PQPQPQ PHLDA1 MYST3 (histone acetyltransferase)	6
CPEB2 (cytoplasmic polyadenylation element binding	
protein) PYPQPP SHISA5	6
Q6 LMX1A NOS1AP	6
QILQQQ FOXP2	6
QPQPQP MEF2A (myocyte enhancer factor) MED12L	6
(Mediator complex)	0
Q-QGSV GRIP1(glutamate receptor interacting	6
protein)	
QQLCCQ SNX6 (sorting nexin)	6
QQPPQQ SMARCA2	6
QQPQQQ SP4	6
Q-QQLP DTNBP1 (Dysbindin)	6
QSGV-P MAP6	6
Q-SLQQ SLC1A6 (glutamate transporter) SQVS	6
ABCA13 GABRA6 (GABA receptor) QUAQGS SLC25A14 (mitochondrial carrier, brain)	6
	6
SSQVSL PIK3C2G (phosphatidylinositol 3-kinase) VQPQQ STRN	6
LQLQP DISC1	6 5
LULUF DIOUT	5

Table 3

The number of gliadin matches (tetrapeptide or more) within human proteins

GENE SYMBOL	Ν
MACF1	24
ATP2A2	21
ANK3	19
RAI1	17
KCNN3, SMARCA2	15
SETD2, SP4, STAB1	14
ABCA13, NUMBL, SYN3	13
ADAM12, ATF7IP	12
ASTN1, BTN3A2, KIF13A, MED25,PCNT, PLXNA2, SPTAN1	11
CCDC141, CIT, FOXP2, RIMS2	10
ABCA7, CABIN1, MAGEL2, MAP1A, MAP6,MYCBP2, PPP1R3F	9
ATF5, EPB41L1, MYO1D, NR4A3, PCM1, RELN	8
CLINT1, HDAC9, ITIH4, KCNH5, LMX1B,	7
MMS22L, SELIL3, SEMA3D, SPTNB4, SYN2, UTRN	7
DNAJC6, HDAC10, LMX1A, POM121L2, RAPGEF6, SIM1, SMARCC1	6
ANK2, BAP1, BRD1, CACNA1B, CDC42SE2, CELF4, DISC1, EGR4, FNIP1, FUBP1, GPR125, GRIP1, HIPK3, MDGA1, NALCN, NCS1, NOTCH4,, TMEM200C	5

Fig 1: The sequence of the gliadin protein used in this study: Polyglutamine or other repeat sequences are highlighted in bold or in grey.

Fig 2:

Top:

The B cell epitope index values of the gliadin protein (amino acid 1-276) as predicted by the Bepipred server. The cut-off value of 0.35 is illustrated by the thick horizontal line.

Bottom

The B cell epitope indices for various glutamine repeats (QQQ) * 1-7

Fig 3

An interactome network for the proteins identified in this study. Proteins in grey boxes show a significant degree of homology with gliadin in the BLAST analyses, or are polyglutamine repeat proteins. Linked boxes represent binding between two components; these data were gleaned from the protein interaction section of Entrez Gene. Arrows indicate other effects on transcription or other functional effects. See Table 1 for details.

Fig 4

A) Clustal line up between gliadin and the potassium channel, KCNN3

B) Gliadin consensus sequences with the KCNN3 protein

C) KCNN3 consensus sequences within the gliadin protein

Fig 5

The B cell epitope index for KCNN3 sequences within the gliadin protein. Other human proteins trapped in passing are also shown.

Fig 6

The B cell epitope indices for sequences of glutamate (top) and myelin related proteins (bottom) sharing consensus sequences with gliadin.

Fig 7

The B cell epitope indices for sequences of schizophrenia-related autoantigens sharing consensus sequences with gliadin.

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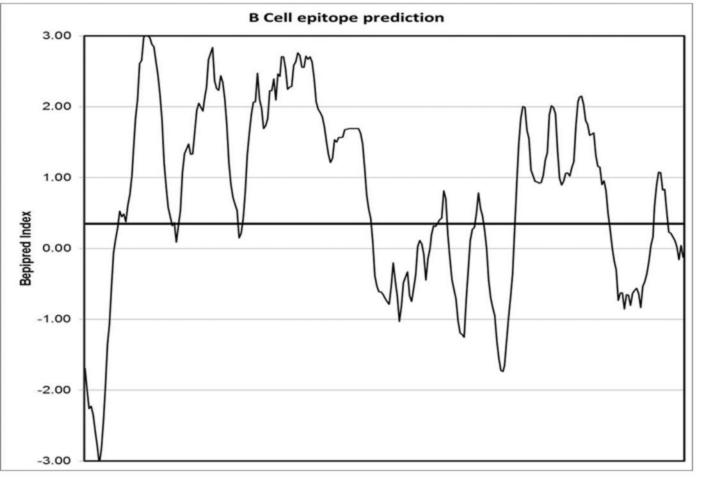
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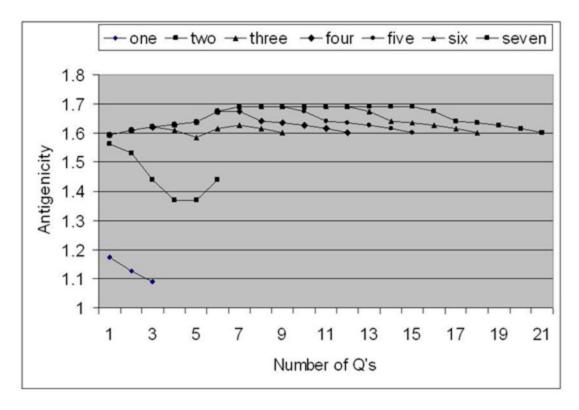
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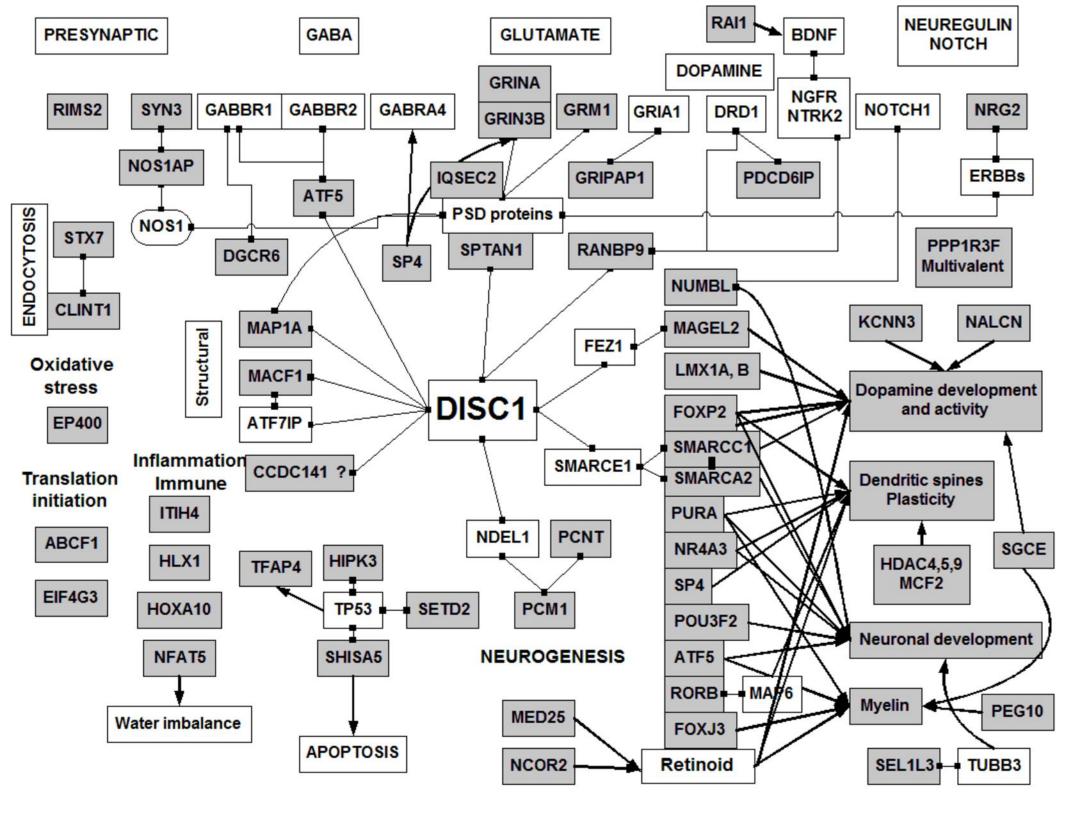
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>gi|100783|pir||A27319 gliadin - wheat Q =glutamine







Gliadin	MKTFLILALLAIVATTATTAVRVPVPQPQPQNPSQPQPQRQVPLVQQQQFPGQQQQFP	58
KCNN3	$\tt MDTSGHFHDSGVGDLDEDPKCPCPSSGDEQQQQQQQQQQQPPPPAPPAAPQQPLGPSLQ$	60
	. : .: *. : *: .* * *:* * * * .:	
Gliadin	PQQPYPQPQPFPSQQPYLQLQPFPQPQPFPPQLPYPQPPFSPQQPYPQPQPQ	111
KCNN3	PQPPQLQQQQQQQQQQQQQPPHPLSQLAQLQSQPVHPGLLHSSPTAFRAPPSSNSTAIL	120
	** * * * * * * * * * * * * * * * * * * *	
Gliadin	YPQPQQPISQQQAQQQQQQQQQQQQQQQILPQILQQQLIPCRDVVLQQHN-	164
KCNN3	HPSSRQGSQLNLNDHLLGHSPSSTATSGPGGGSRHRQASPLVHRRDSNPFTEIAMSSCKY	180
	:*:* . : : : : :::* * : ::: * :: :	
Gliadin	IAHARSQVLQQSTYQPLQQLCCQQLWQIPEQSRCQAIHNVVHAIIL	210
K C N N 3	$\verb+SGGVMKPLSRLSASRRNLIEAETEGQPLQLFSPSNPPEIVISSREDNHAHQTLLHHPNAT$	240
	* *. : : **** : .: :* .** : *.	
GliadinHQ	QQQQQQPSSQVSLQQPQQQYPSGQGFFQPSQQNPQAQG250	J
K C N N 3	HNHQHAGTTASSTTFPKANKRKNQNIGYKLGHRRALFEKRKRLSDYALIFGMFGIVVMVI	300
::: .		
Gliadin-S	SVQPQQLPQFEEIRNLALQTLPRMCNVYIPPYCSTTTAPFGIFGTN 296	
KCNN3 E	TELSWGLYSKDSMFSLALKCLISLSTIILLGLIIAYHTREVQLFVIDNGADDWRIAMTY 36	0
: . * .		

Identities = 54/300 = 18%

B)Tetrapeptide or greater gliadin consensus sequences within the KCNN3 protein

>gi|116805330|ref|NP_002240.3| small conductance calcium-activated potassium channel protein 3 isoform a [Homo sapiens]

Identities= 38/720 5.3%

C)KCNN3 matching contiguous sequences within the gliadin protein

>gi:100783:pir::A27319 gliadin - wheat

Identities 94/296=32%

a

		Bepipred	ed index		
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QPQP ABCA7 MED12L ME AED12L ME AAGEL2 MEF2A QPQPQ ALX KC QQQQ ABCF1 F AQQQ ABCF1 F ACOR2 NFAT5 AQQQ ABCF1 F ACOR2 NFAT5 PQQP ABCA7 ALX4 MED12L MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 MED12L MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ABCA7 ALX4 MED12L MED1 PQQP ABCA7 ALX4 MED12L MED1 PQQP ABCA7 ALX4 ACT III MED1 PQQP ACT III MED1 ACT III MED1 PQQP AC	QPQP ABCA7 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 QPQPQ ALX4 [QPQP] ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL QQQQ ABCF1 ARID3B ATRX FOXP2 GRIN3A GRM1 HDAC5 HDAC9 HLX KCNN3 LMX1A LMX1B MAP1A NCOR1 NCOR2 QQQQ ABCF1 FOXP2 HDAC9 KCNN3 LMX1A LMX1B MAP1A NCOR2 NFAT5 KCNN3 LMX1A LMX1B MAP1A NCOR1 NCOR2 PQQA AFF5 CLINT1 EPB41L1 IQSEC2 KCNN3 KIF13A NCOR1 NCOR2 NEAT5 NUMBL PCM1 DNCOR1 NCOR2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 AGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 KF3A MC02 KCNN3 KF3A MC02 KCNN3 KF3A KCNN3 KF3A MC02 KCNN3 KF3A MC02 KCNN3 KF3A KCNN3 KF	 QPQP ABCA7 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 QPQPQ ALX4 [QPQP] ABCA7 ALX4 CDC42SE2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL QQQQ ABCF1 ARID3B ATRX FOXP2 GRIN3A GRM1 HDAC5 HDAC9 HLX KCNN3 LMX1A LMX1B MAP1A NCOR1. NCOR2 PPQQQ ABCF1 FOXP2 HDAC9 KCNN3 LMX1A LMX1B MAP1A NCOR2 NFAT5 NUMBL PCM1 PHC2 POU3F2 RAI1 RIMS2 PPQQP SYN2 [PPQQP] ATF5 KCNN3 RAI1 [PPQQ] BAP1 BRPF1 FIXP3 GPR125 LMX1B MAP6 MED25 PQQQ ABC71 EPB41L1 IQSEC2 KCNN3 KIF13A NCOR1 NCOR2 NFAT5 NUMBL PCM1 DNCOR1 SYN2 PPQQP ATF5 CLINT1 EPB41L1 IQSEC2 KCNN3 MGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PPQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PPQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 MAGEL2 MEF2A MED12L MED15 MMS22L MYO1D NUMBL RAI1 SETD2 PQQP ABCA7 ALX4 CDC425E2 DNAJC6 KCNN3 KIF13A NCOR1 NCOR2 SEMA3D SYN2 QQQUI CHERP CKB [QQQL] ASTN2 QQQUI CHERP CKB [QQQL] ASTN2 QQQUI CH			

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