Supplementary Table 1: Primers used for genotyping, cloning of DBH promoter region and composition of PCR used for Genotyping DBH polymorphisms

Marker	Oligo	Sequence	Allele	Product Size (bp)	Restriction enzyme	Fragment size (bp)
rs141116007	19bpDBHF	5'-AGGACCCACCACCATCCT-3'	Ins	202		202
	19bpDBHR	5'-CCTCAATCTTGGGGCTGA-3'	Del			184
rs1611115	rs1611115F	5'-CAGACCTGCTGTCATGGGTA -3'	C	476	Fau I	254+222
	rs1611115R	5'-AAGGAAAAGAGGGTGAGTGACA -3'	T		(New England Biolabs, Ipswich, MA, USA)	476
rs1989787	rs1989787F	5'-TCTCTCCCCTCTGGGGTATT-3'	A	445	Eco47III (AfeI)	445
	rs1989787R	5'-GCTAGAGGAAAGCCAGTTTTGA-3'	G		(Thermo Fisher Scientific, Waltham, MA,USA)	329+116

Composition of PCR used to genotype DBH markers

	rs141116007	rs1989787	rs1611115
Components			
10x Reaction buffer MgCl ₂ free	2.0	2.0	1.0
50mM MgCl ₂	0.6	0.8	0.4
2.5mM dNTPS	1.6	2.0	1.0
Forward Primer (10pmoles/µl)	1.0	1.0	0.25
Reverse Primer (10pmoles/µl)	1.0	1.0	0.25
3B DNA polymerase (5U/µl)	0.5	0.2	0.1
H_2O	11.3	11.0	6.0
DNA (25ng/μl)	2.0	2.0	1.0
Reaction volume(µl)	20	20	10

Primers used for cloning DBH promoter region and thermal cycling conditions for the PCR

Name	Sequence [#]	Tm°∗	Amplicon length	

DBH 19bp incF	5'-CTATACGCGTTCTGTTCCCACCTTTTCCAG-3'	72.1/58.4	5677bp
Prom 2	5'-GCTTCTCGAGGGCTGGGGTGAGCTCTCT-3'	76/60.7	
DelXIX F	5'-CTGTCTCGCTAGCCCTCACTGCAT-3'	68.7	2262bp
DelXIX R	5'-TTGGCCACATGCCTCCCTGT-3'	62.5	

#Sequence from 5' to the end of highlighted region are overhangs and highlighted region is restriction sites (Mlu I and Xho I respectively) followed by sequence complimentary to genomic DNA

Supplementary Table 2: PCR cycling, restriction digestion conditions for genotyping and thermal cycling conditions for amplicons used in cloning DBH promoter region

Steps	Thermal	cycling condition fo	or rs141116007 amplicon	Thermal o	ycling condition for	rs1989787 amplicon	Thermal o	ycling condition for	rs1611115 amplicon
	°C	Time	Cycles	°C	Time	Cycles	°C	Time	Cycles
1	95	5min		94	5min		94	5min	_
2	95	15s	Step 2 to 4	94	1min	Step 2 to 4	94	1min	Step 2 to 4
3	59	30s	35 cycles	58	30s	40 cycles	58	30s	35 cycles
4	72	30s		72	1min		72	1min	
5	72	7min		72	15min		72	15min	
7	4	∞		4	∞		4	∞	
			Destriction	digastian car	ditions for re102079	7 and re1611115			

		Restricti	tion digestion conditions for rs1989/8/ and rs1611115
rs19897	187	rs161	11115
Components	Volume (µl)	Components	Volume (μl)
10 x Buffer O	2.5	10 x Cutsmart™	1.5
		buffer	
PCR reaction	20.0	PCR reaction	10.0
H_2O	2.4	H_2O	3.3
Eco47III (AfeI) 10 U/ μl	0.1	FauI 5U/ μl	0.2
Incubate 37°C O/N		Incubate 55°C O/N	

Steps	Cycling cond	lition for Insert I		Cycling con	ditions for Insert I	I	
1	94°C	1min		94°C	1min		
2	98°C	10s	Step 2 to 4	98°C	10s	Step 2 to 4	
3	62°C	30s	35 cycles	66°C	30s	35 cycles	
4	72°C	5.6min	-	72°C	2.2min	•	
5	72°C	10min		72°C	10min		
6	4°C	∞		4°C	∞		

^{*}Tm of the entire oligo / Tm of the omplimentary region

Supplementary Table 3: Associations of DBH markers with accuracy of different cognitive domains

	Abstraction and Mental flexibility _{accuracy}	Attention _{accuracy}	Face memory _{accuracy}	Spatial memory _{accuracy}	Verbal memory _{accuracy}	Spatial processing _{accuracy}	Sensorimotor dexterity accuracy	Emotional processing accuracy
Gender	F(1,608)=13.3;p<0.001						F(1,586)=13.4;p<0.001	
Health	F(1,608)=50.3;p<0.001	F(1,529)=20.7;p<0.00	F(1,608)=17.3;p<0.001	F(1,605)=7.6;p=0.006	F(1,583)=33.4;p<0.001	F(1,515)=6.1;p<0.0 5	F(1,586)=15.1;p<0.001	F(1,603)=15.4;p<0.001
Indel19bp								
rs1989787								
rs1611115								
Health * Indel19bp								
Health * rs1989787	F(2,608)=3.7;p=0.03							
Health * rs1611115					F(2,583)=3.1;p=0.048			F(2,603)=3.9;p=0.05
Indel19bp * rs1989787								
Indel19bp * rs1611115								
_								
rs1989787 * rs1611115 Health * Indel19bp *	F(3,608)=3.0;p=0.03							
rs1989787								
Health * Indel19bp *	F(4,608)=3.3;p=0.01							
rs1611115								
Health * rs1989787 *		F(1,529)=4.5;p=0.03						F(1,603)=4.4;p=0.04
rs1611115								
Indel19bp * rs1989787 *								
rs1611115								
Health * Indel19bp *	F(2,608)=4.1;p=0.02							
rs1989787 * rs1611115								

Supplementary Table 4: Associations of DBH markers with processing speed of different cognitive domains

	Abstraction and Mental	Attention _{processing} speed	Face memory _{processing}	Spatial memory _{processing}	Verbal	Spatial Processing speed	Sensorimotor dexterity _{processing speed}	Emotional processing speed
	flexibility _{processing speed}		speed	speed	memory _{processing} speed	Sprocessing speed	processing speed	F- 3 3 3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
Gender		F(1,527)=13.9;p<0.001					F(1,582)=18.3;p<0.001	F(1,598)=6.1;p<0.05
Health	F(1,601)=6.2;p<0.05	F(1,527)=27.1;p<0.001	F(1,603)=4.0;p=0.05		F(1,578)=3.8;p=0.05	F(1,504)=17.1;p<0.001	F(1,582)=12.6;p<0.001	F(1,598)=17.0;p<0.001
Indel19bp								
rs1989787								
rs1611115								
Health * Indel19bp								F(2,598)=3.4;p=0.03
Health * rs1989787								
Health * rs1611115								
Indel19bp * rs1989787	F(3,601)=2.9;p=0.03							
Indel19bp * rs1611115								
rs1989787 * rs1611115								
Health * Indel19bp *								
rs1989787								
Health * Indel19bp *								
rs1611115								
Health * rs1989787 *								
rs1611115								
Indel19bp * rs1989787 *	F(2,601)=5.1;p=0.006							
rs1611115								
Health * Indel19bp *								
rs1989787 * rs1611115								

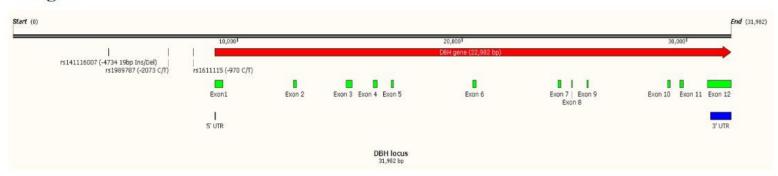
Supplementary Table 5: Associations of DBH markers with efficiency of different cognitive domains

	Abstraction and Mental flexibility _{efficiency}	Attention _{efficiency}	Face memory _{efficiency}	Spatial memory _{efficiency}	Verbal memory _{efficiency}	Spatial Processing _{efficiency}	Sensorimotor dexterity _{efficiency}	Emotional processing _{efficiency}
Gender	F(1,601)=14.2;p<0.001						F(1,582)=14.5;p<0.001	
Health	F(1,601)=50.7;p<0.001	F(1,527)=25.0;p<0.001	F(1,603)=18.3;p<0.001	F(1,600)=6.9;p=0.009	F(1,578)=29.4;p<0.001	F(1,504)=8.6;p<0.003	F(1,582)=17.4;p<0.001	F(1,593)=16.7;p<0.001
Indel19bp								
rs1989787								
rs1611115								
Health * Indel19bp								
Health * rs1989787	F(2,601)=3.5;p=0.031							
Health * rs1611115								
Indel19bp * rs1989787								
Indel19bp * rs1611115								
rs1989787 * rs1611115								
Health * Indel19bp *	F(3,601)=2.9;p=0.03							
rs1989787								
Health * Indel19bp *	F(4,601)=3.1;p=0.02							
rs1611115								
Health * rs1989787 *		F(1,527)=4.1;p=0.04						
rs1611115								
Indel19bp * rs1989787 *								
rs1611115								
Health * Indel19bp *	F(2,601)=4.4;p=0.01							
rs1989787 * rs1611115								

Supplementary Figure 1: Position of *DBH* variants selected for genotyping with respect to the gene

The three variants that were studied 19bp *Ins/Del*, rs1989787 and rs1611115 are marked with respective to DBH gene. The Exons are drawn to scale. Linkage disequilibrium (LD) measures (D' and R-sq) between the three variants are given in the table

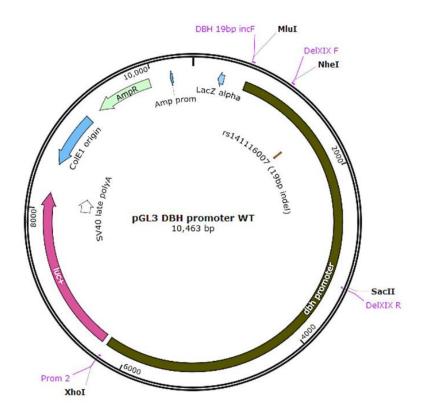
Supplementary Figure 1: Position of *DBH* variants selected for genotyping with respect to the gene



Marker 1	Marker 2	D'	R-Sq
rs141116007	rs1611115	0.23	0.02
rs141116007	rs1989787	0.57	0.05
rs1989787	rs1611115	0.82	0.03

Supplementary Figure 2: DBH WT promoter construct used for functional analysis
pGL3 DBH promoter WT construct (10.5kb) with 19bp <i>Ins</i> allele(rs141116007) with the primers used for cloning namely DBH 19bp incF and prom2 for amplicon I with Mlu I and Xho I overhangs; Del XIXF and Del XIXR for amplicon II with Nhe I and Sac II are also depicted

Supplementary Figure 2: DBH WT promoter construct used for functional analysis



Supplementary Figure 3: A decrease in accuracy, processing speed and efficiency of PennCNB neurocognitive scores in schizophrenia subjects observed with Cluster analysis

A general decrease in all the 24 cognitive scores was observed in SZ subjects as compared to healthy controls. Accuracy, processing speed and efficiency of cognitive scores of abstraction and flexibility, attention, working memory, face memory, spatial memory, spatial processing, sensorimotor dexterity and emotional processing are illustrated.

Supplementary Figure 3: A decrease in accuracy, processing speed and efficiency of PennCNB neurocognitive scores in schizophrenia subjects observed with Cluster analysis

