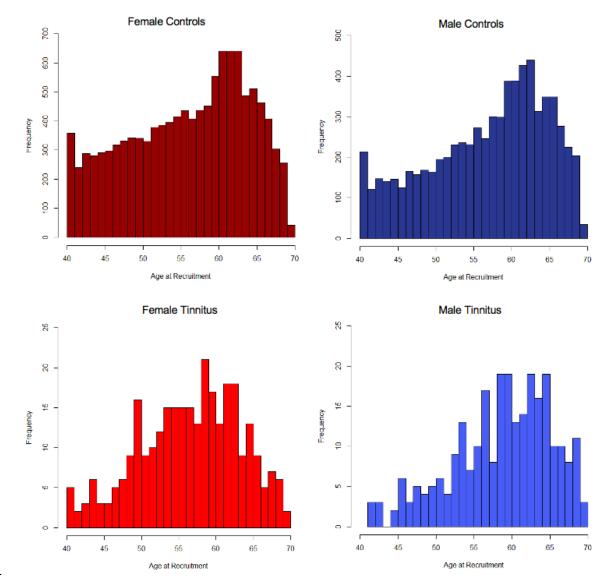
493 Genetic predisposition to tinnitus in the UK Biobank population

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500 Supplementary Information

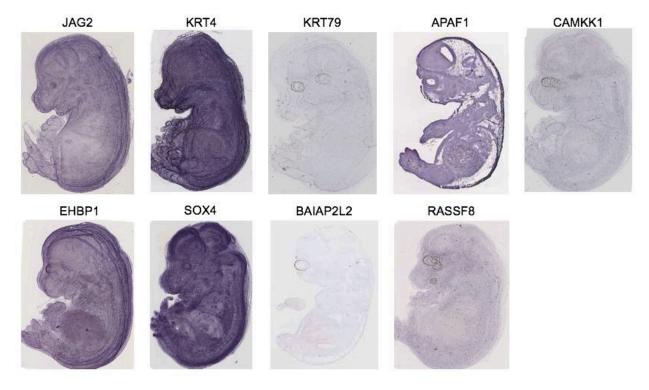


Fig S1. Age distribution of all included participants in both tinnitus and control cohorts. At the
 time of recruitment, all participants fell between 40 and 70 years of age. In all four groups,
 distribution was approximately equivalent, with the peak of participants being 60-65 years old.

	Field Subject	Field Number	Categories Excluded
Ethnicity			
	Genetic ethnic grouping	22006	Non-Caucasian
Environment			
	Noisy workplace	4825	Yes for ≥ 1 year
	Loud music exposure	4836	Yes for ≥ 1 year
lle enire e		4030	
Hearing		00.17	
	Hearing difficulty	2247	Yes or total deafness
	Hearing aid	3393	Yes
	Cochlear implant	4792	Yes Death, hospital admissions, primary
	Unsuppurative otitis media	131229	care records
	Suppurative otitis media	131231	Death, hospital admissions, primary care records
			Death, hospital admissions, primary
	Eustachian obstruction	131235	care records Death, hospital admissions, primary
	Mastoiditis	131239	care records Death, hospital admissions, primary
	Choleseatoma of middle ear	131241	care records
	Perforation of tympanic membrane	131243	Death, hospital admissions, primary care records
			Death, hospital admissions, primary
	Otosclerosis Conductive and sensorineural hearing	131251	care records Death, hospital admissions, primary
	loss	131259	care records
	Otalgia and effusion of ear	131263	Death, hospital admissions, primary care records
Cardiovascular			
	Stroke	42006	Death, hospital admissions, primary care records
		42000	Death, hospital admissions, primary
Central Nervous	Hypertensive disease	131289	care records
System			
	Type of cancer	40006	Malignant neoplasms of eye, brain, and other parts of CNS
		120002 120005	Death, hospital admissions, primary
	Bacterial meningitis	130993, 130995	care records Death, hospital admissions, primary
	Meningitis Encephalitis, myelitis, and	130997, 130999	care records Death, hospital admissions, primary
	encephalomyelitis	131001, 131003	care records
	Intracranial and intraspinal abscess and granuloma	131005, 131007	Death, hospital admissions, primary care records
	Intracranial and intraspinal phlebitis and		Death, hospital admissions, primary
	thrombophlebitis	131009	care records Death, hospital admissions, primary
	Huntington's disease	131013	care records Death, hospital admissions, primary
	Hereditary ataxia	131015	care records
	Spinal muscular atrophy	131017	Death, hospital admissions, primary care records
			Death, hospital admissions, primary
	Postpolio syndrome	131021	care records Death, hospital admissions, primary
	Parkinson's disease	131023	care records

		Death, hospital admissions, primary
Secondary Parkinsoni	sm 131025	
		Death, hospital admissions, primary
Dystonia	131031	
		Death, hospital admissions, primary
Alzheimer's disease	131037	
		Death, hospital admissions, primary
Multiple sclerosis	131043	
		Death, hospital admissions, primary
Epilepsy	131049	care records
		Death, hospital admissions, primary
Status epilepticus	131051	care records
		Death, hospital admissions, primary
Transient cerebral isc	haemic attacks 131057	care records
Vascular syndromes of	of the brain in	Death, hospital admissions, primary
verebrovascular disea	se 131059	e care records
		Death, hospital admissions, primary
Disorders of the trigen	ninal nerve 131063	3 care records
		Death, hospital admissions, primary
Facial nerve disorders	131065	5 care records
		Death, hospital admissions, primary
Nerve root and plexus	disorders 131071	care records
Myasthenia gravis and	d other myoneural	Death, hospital admissions, primary
disorders	131093	3 care records
		Death, hospital admissions, primary
Infantile cerebral pals	/ 131101	care records
		Death, hospital admissions, primary
Hydrocephalus	131111	care records
		Death, hospital admissions, primary
Toxic encephalopathy	131113	3 care records

 Table S1. UK Biobank fields disqualified from GWAS.



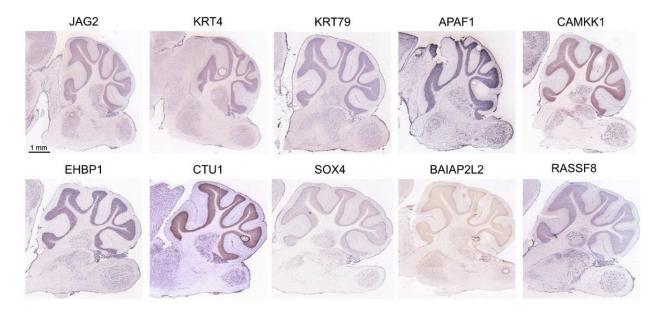
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509 **Fig S2.** *In situ* hybridization of additional gene hits in 14.5 embryonic mice. Genes already

available on the eMouseAtlas exhibit varying levels of expression in the brain during this stage

of development, with *Baiap2l2* and *Krt79* showing minimal expression. Image credit: EMAGE

512 Project (EMAGE: 3758, 16846, 14151, 12754, 14122, 12668, 13766, 29351, 11832).



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- 514 **Fig S3**. *In situ* hybridization of additional gene hits in adult mice. All genes already
- recorded by the Allen Brain Atlas show some level of mRNA expression, with *Camkk1*, *Ctu1*,
- and *Rassf8* showing the most elevated levels across samples. Scaling is equal across images.
- 517 Image credit: Allen Institute (https://mouse.brain-map.org/experiment/show?id=69735652,
- 518 https://mouse.brain-map.org/experiment/show?id=69735956, https://mouse.brain-
- 519 map.org/experiment/show?id=68522308, https://mouse.brain-
- 520 map.org/experiment/show?id=68745275, https://mouse.brain-
- 521 map.org/experiment/show?id=70431211, https://mouse.brain-
- 522 map.org/experiment/show?id=70239099, https://mouse.brain-
- 523 map.org/experiment/show?id=413826, https://mouse.brain-
- 524 map.org/experiment/show?id=81654151, https://mouse.brain-
- 525 map.org/experiment/show?id=74431104, https://mouse.brain-
- 526 map.org/experiment/show?id=69443280).

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