

the first symptom of a serious condition such as a vestibular schwannoma. Pulsatile tinnitus requires evaluation of the vascular system of the head to exclude vascular tumors such as glomus jugulare tumors, aneurysms, dural arteriovenous fistulas, and stenotic arterial lesions; it may also occur with SOM, superior semicircular dehiscence, and inner-ear dehiscence. It is most commonly associated with some abnormality of the jugular bulb such as a large jugular bulb or jugular bulb diverticulum. In absence of demonstrated pathology on MRA/MRV or CT angiography, pulsatile tinnitus is usually attributed to turbulent venous blood flow through the transverse sinus, sigmoid sinus, and the jugular bulb.

■ GENETIC CAUSES OF HEARING LOSS



More than half of childhood hearing impairment is thought to

be hereditary; hereditary hearing impairment (HHI) can also manifest later in life. HHI may be classified as either nonsyndromic, when hearing loss is the only clinical abnormality, or syndromic, when hearing loss is associated with anomalies in other organ systems. Nearly two-thirds of HHIs are nonsyndromic. Between 70% and 80% of nonsyndromic HHI is inherited in an autosomal recessive manner and designated DFNB; another 15–20% is autosomal dominant (DFNA). Less than 5% is X-linked (DFNX) or maternally inherited via the mitochondria.

More than 150 loci harboring genes for nonsyndromic HHI have been mapped, with recessive loci outnumbering dominant ones; numerous genes have now been identified ([Table 34-1](#)). The hearing genes fall into the categories of structural proteins (*MYH9*, *MYO7A*, *MYO15*, *TECTA*, *DIAPH1*), transcription factors (*POU3F4*, *POU4F3*), ion channels (*KCNQ4*, *SLC26A4*), and gap junction proteins (*GJB2*, *GJB3*, *GJB6*). Several of these genes, including *GJB2*, *TECTA*, and *TMC1*, cause both autosomal dominant and recessive forms of nonsyndromic HHI. In general, the hearing loss associated with dominant genes has its onset in adolescence or adulthood, varies in