TABLE 82-3 SELECTED EXAMPLES OF LOCUS HETEROGENEITY AND PHENOTYPIC HETEROGENEITY

Phenotypic Heterogeneity

Gene, Protein	Phenotype	Inheritance	OMIM
LMNA, Lamin A/C	Emery-Dreifuss muscular dystrophy (AD)	AD	181350
	Familial partial lipodystrophy Dunnigan	AD	151660
	Hutchinson-Gilford progeria	AD	176670
	Atypical Werner's syndrome	AD	150330
	Dilated cardiomyopathy	AD	115200
	Early-onset atrial fibrillation	AD	607554
	Emery-Dreifuss muscular dystrophy (AR)	AR	604929
	Limb-girdle muscular dystrophy type 1B	AR	159001
	Charcot-Marie-Tooth type 2B1	AR	605588
KRAS	Noonan's syndrome	AD	163950
	Cardio-facio-cutaneous syndrome	AD	115150

Locus Heterogeneity

Phenotype	Gene	Chromosomal Location	Protein
Familial hypertrophic cardiomyopathy	*MYH7*	14q12	Myosin heavy chain beta
Genes encoding sarcomeric proteins	*TNNT2*	1q2	Troponin-T2
	TPM1	15q22.1	Tropomyosin alpha
			Myosin-binding
	MYBPC3	11p11q	protein C
	TNNI3	19q13.4	Troponin 1
	MYL2	12q23-24.3	Myosin light chain 2
	MYL3	3p	Myosin light chain 3
	TTN	2q24.3	Cardiac titin
	ACTC	15q11	Cardiac alpha actin
	MYH6	14q1	Myosin heavy chain alpha
	MYLK2	20q13.3	Myosin light-peptide kinase
	CAV3	3p25	Caveolin 3
Genes encoding nonsarcomeric proteins	*MTT1*	Mitochondrial	tRNA isoleucine
	MTTG	Mitochondrial	tRNA glycine
	PRKAG2	7q35-q36	AMP-activated protein kinase γ2 subunit
	DMPK	19q13.2-13.3	Myotonin protein kinase (myotonic dystrophy)
	FRDA	9q13	Frataxin (Friedreich's ataxia)
Polycystic kidney disease	*PKD1*	16p13.3-13.12	Polycystin 1 (AD)
	PKD2	4q21.-23	Polycystin 2 (AD)
	PKHD1	6p21.1-p12	Fibrocystin (AR)
Noonan's syndrome	*PTPN11*	12q24.1	Protein-tyrosine phosphatase 2c
	KRAS	12p12.1	KRAS

of the helical collagen fiber. Similarly, muscular dystrophy syndromes can be caused by mutations in various genes, consistent with the fact that it can be transmitted in an X-linked (Duchenne or Becker), autosomal dominant (limb-girdle muscular dystrophy type 1), or autosomal recessive (limb-girdle muscular dystrophy type 2) manner (Chap. 462e). Mutations in the X-linked *DMD* gene, which encodes dystrophin, are the most common cause of muscular dystrophy. This feature reflects the large size of the gene as well as the fact that the phenotype is expressed in hemizygous males because they have only a single copy of the X chromosome. Dystrophin is associated with a large protein complex linked to the membrane-associated cytoskeleton in muscle. Mutations in several different components of this protein complex can also cause muscular dystrophy syndromes. Although the phenotypic features of some of these disorders are distinct, the phenotypic spectrum caused by mutations in different genes overlaps, thereby leading to nonallelic heterogeneity. It should be noted that mutations in dystrophin also cause allelic heterogeneity. For example, mutations in the *DMD* gene can cause either Duchenne's or the less severe Becker's muscular dystrophy, depending on the severity of the protein defect.

Recognition of nonallelic heterogeneity is important for several reasons: (1) the ability to identify disease loci in linkage studies is reduced by including patients with similar phenotypes but different genetic disorders; (2) genetic testing is more complex because several different genes need to be considered along with the possibility of different mutations in each of the candidate genes; and (3) novel information is gained about how genes or proteins interact, providing unique insights into molecular physiology.

Phenocopies refer to circumstances in which nongenetic conditions mimic a genetic disorder. For example, features of toxin- or drug-induced neurologic syndromes can resemble those seen in Huntington's disease, and vascular causes of dementia share phenotypic features with familial forms of Alzheimer's dementia (Chap. 448). As in nonallelic heterogeneity, the presence of phenocopies has the potential to confound linkage studies and genetic testing. Patient history and subtle differences in phenotype can often provide clues that distinguish these disorders from related genetic conditions.

VARIABLE EXPRESSIVITY AND INCOMPLETE PENETRANCE The same genetic mutation may be associated with a phenotypic spectrum in different affected individuals, thereby illustrating the phenomenon of *variable expressivity*. This may include different manifestations of a disorder variably involving different organs