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**Lab Dept:** Chemistry

**Test Name:** ALPHA-1-ANTITRYPSIN PHENOTYPE

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***General Information***

**Lab Order Codes:** A1AP

**Synonyms:** A1A Phenotype, serum; AAT Phenotype, serum; Pi Typing; Protease Inhibitor Allo Typing

**CPT Codes:** 82103 - Alpha-1-antitrypsin; total  
82104 - Alpha-1-antitrypsin; phenotype

**Test Includes:** Alpha-1-Antitrypsin, serum reported in mg/dL and phenotype.  
**NOTE: DO NOT order with Alpha-1-Antitrypsin Proteotype Deficiency Panel.** For more information on this testing, see [Mayo's Testing Algorithm](#)

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***Logistics***

**Test Indications:** Useful for identification of homozygous and heterozygous phenotypes of alpha-1-antitrypsin deficiency.

**Lab Testing Sections:** Chemistry - Sendouts

**Referred to:** Mayo Medical Laboratories (MML Test: A1APP)

**Phone Numbers:** MIN Lab: 612-813-6280  
STP Lab: 651-220-6550

**Test Availability:** Daily, 24 hours

**Turnaround Time:** 2 - 6 days, performed Monday - Friday

**Special Instructions:** N/A

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***Specimen***

**Specimen Type:** Blood

**Container:** SST (Marble, gold or red)

**Draw Volume:** 4 mL (Minimum: 1.5 mL) blood

**Processed Volume:** 1.25 mL (Minimum: 0.5 mL) serum

<b>Collection:</b>	Routine venipuncture
<b>Special Processing:</b>	Lab Staff: Centrifuge specimen, remove serum aliquot into a screw-capped round bottom plastic vial. Store and ship at refrigerated temperatures. Forward promptly.
<b>Patient Preparation:</b>	None
<b>Sample Rejection:</b>	Specimens other than serum; gross lipemia; mislabeled or unlabeled specimens

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### ***Interpretive***

**Reference Range:** **Alpha-1-Antitrypsin, Serum:** 100 - 190 mg/dL

**Alpha-1-Antitrypsin Phenotyping:** The interpretive report will identify the alleles present. For rare alleles, the report will indicate whether or not they have been associated with reduced quantitative levels of alpha-1-antitrypsin.

Most normal individuals have the M phenotype (M, M1 or M2). Over 99% of M phenotypes are genotypically MM. In the absence of family studies, the phenotype (M) and quantitative level can be used to infer the genotype (MM). The most common alleles associated with a quantitative deficiency are Z and S. The reports for the rare alleles will indicate whether or not they have been associated with reduced quantitative levels.

Interpretation: There are >40 Alpha-1-antitrypsin (A1A) phenotypes (most of these are associated with normal quantitative levels of protein). The most common normal phenotype is M (M, M1 or M2), and >90% of Caucasians are homozygous M (MM) genotype.

A1A deficiency is usually associated with the Z phenotype (ZZ genotype), but genotypes such as SS and SZ are also associated with decreased A1A levels.

**Critical Values:** N/A

**Limitations:** This assay identifies the phenotype of the circulating alpha-1-antitrypsin (A1A) protein. If the patient is already on replacement therapy, the phenotype will detect patient and replacement protein.

If two bands are seen, such as an M band and a Z band, it is reported as MZ (eg, heterozygous).

If one band is seen, such as the Z band and the quantitative level is consistent with a homozygote, the phenotype is assumed to be homozygous and is reported as ZZ.

**Methodology:** Isoelectric Focusing and Nephelometry

**References:**

[Mayo Medical Laboratories Web Page](#) (August 2016)