

Names for RHAG (ISBT 030) Blood Group Alleles

General description: The RHAG blood group system consists of three antigens carried on a multipass membrane glycoprotein called RhAG (Rh-associated glycoprotein; aka CD241). It consists of 409 amino acids and both amino and carboxyl termini are predicted to be intracellular.

Gene name: *RHAG*
 Number of exons: 10
 Initiation codon: Beginning of exon 1
 Stop codon: End of exon 10
 Entrez Gene ID: 6005
 LRG Sequence: NG_011704.1 (genomic)
 NM_000324.2 (transcript)
 Reference allele: *RHAG*01* (shaded)

| Reference allele <i>RHAG*01</i> encodes RHAG1, RHAG3 | | | | |
|--|--------------------|-----------------------|----------|-----------------------------|
| Phenotype | Allele name | Nucleotide change | Exon | Predicted amino acid change |
| RHAG:1 or Duclos+ | <i>RHAG*01</i> | | | |
| RHAG:-1 or Duclos- | <i>RHAG*-01</i> | c.316C>G | 2 | p.Gln106Glu |
| RHAG:2 or OI(a+) | <i>RHAG*02</i> | c.680C>T | 5 | p.Ser227Leu |
| RHAG:-3 or DSLK- | <i>RHAG*-03</i> | c.490A>C | 3 | p.Lys164Gln |
| RHAG:4 | <i>RHAG*04</i> | c.808G>A | 6 | p.Val270Ile |
| Null phenotypes | | | | |
| Rh _{null} | <i>RHAG*01N.01</i> | c.154_157delinsGA | 2 | p.Pro52Aspfs*57 |
| Rh _{null} | <i>RHAG*01N.02</i> | c.1086delA | 8 | p.Ala363Leufs*15 |
| Rh _{null} | <i>RHAG*01N.03</i> | c.157+1G>A | Intron 1 | Alternative splicing |
| Rh _{null} | <i>RHAG*01N.04</i> | c.945+1G>A | Intron 6 | Alternative splicing [1] |
| Rh _{null} | <i>RHAG*01N.05</i> | c.946-1G>A | Intron 6 | Alternative splicing |
| Rh _{null} | <i>RHAG*01N.06</i> | c.946-1G>T | Intron 6 | Alternative splicing |
| Rh _{null} | <i>RHAG*01N.07</i> | c.1067+1G>A | Intron 7 | Alternative splicing |
| Rh _{null} | <i>RHAG*01N.08</i> | c.808G>A; c.838G>A | 6 | p.Val270Ile; p.Gly280Arg |
| Rh _{null} | <i>RHAG*01N.09</i> | c.836G>A | 6 | p.Gly279Glu |
| Rh _{null} | <i>RHAG*01N.10</i> | c.1094T>G | 8 | p.Leu365Arg [1] |

Names for *RHAG* (ISBT 030) blood group alleles v4.0 160621

| | | | | |
|--------------------|--------------------|---------------------------------------|----------|--------------------------------------|
| Rh _{null} | <i>RHAG*01N.11</i> | c.1139G>T | 9 | p.Gly380Val; Alternative splicing |
| Rh _{null} | <i>RHAG*01N.12</i> | c.353C>T | 3 | p.Ala118Glu [5] |
| Rh _{null} | <i>RHAG*01N.13</i> | c.1003G>A | 7 | p.Gly335Ser |
| Rh _{null} | <i>RHAG*01N.14</i> | c.946-2A>G | Intron 6 | Alternative splicing [7] |
| Rh _{null} | <i>RHAG*01N.15</i> | c.(?-62)_(*638_?)del Gene deletion | 1-10 | p.0 [8] |
| Rh _{null} | <i>RHAG*01N.16</i> | c.310C>T | 2 | p.Gln104Ter [9] |
| Mod phenotypes | | | | |
| Rh _{mod} | <i>RHAG*01M.01</i> | c.1183delA | 9 | p.Asn395Thrfs*68 |
| Rh _{mod} | <i>RHAG*01M.02</i> | c.3G>T | 1 | p.Arg2_Met8del |
| Rh _{mod} | <i>RHAG*01M.03</i> | c.236G>A | 2 | p.Ser79Asn |
| Rh _{mod} | <i>RHAG*01M.04</i> | c.269G>T | 2 | p.Gly90Val [2] |
| Rh _{mod} | <i>RHAG*01M.05</i> | c.398T>C | 3 | p.Leu133Pro [3] |
| Rh _{mod} | <i>RHAG*01M.06</i> | c.560G>A | 4 | p.Gly187Asp [2] |
| Rh _{mod} | <i>RHAG*01M.07</i> | c.1195G>T | 9 | p.Asp399Tyr |
| Rh _{mod} | <i>RHAG*01M.08</i> | c.182T>G | 2 | p.Ile61Arg |
| Rh _{mod} | <i>RHAG*01M.09</i> | c.194T>C | 2 | p.Phe65Ser |
| Rh _{mod} | <i>RHAG*01M.10</i> | c.572G>A | 4 | p.Arg191Gln [6] |

RHAG3 assigned provisionally.

Assignment of null (*N*) and mod (*M*) alleles has been made according to the phenotypic expression of RhD and RhCE antigens.

1. Tsuneyama H, et al. *Transfusion* 2005;45(suppl):130A (abstract).
2. Scharberg A, et al. *Vox Sang* 2006;91(suppl 3):129 (abstract).
3. Tsuneyama H, et al. *Transfusion* 2008;48(suppl):194A-195A (abstract).
4. Poole J, et al. A novel RHAG blood group antigen associated with severe HDFN. *Vox Sang* 2011;101 (Suppl. 1):70 (abstract).
5. Grimsley S, Poole J, Thornton N, et al. Novel mutations in RHAG causing two new examples of the regulator type of Rhnull. *Transfus Med* 2012;22(Suppl.1):21 (abstract).
6. Tsuneyama H, Isa K, Ogasawara K, Yabe R, Uchikawa M, Minami M. Identification of a mutation in the *RHAG* gene of Japanese with weak D phenotype. *Transfusion* 2014;53 (Suppl.):167A (abstract).
7. Arsenovic MG, et al. *Vox Sang* 2014;107(suppl.1):192 (abstract).
8. Gómez-Torreiro, et al. *Transfusion* 2015;55:197-198.
9. Arnoni CP, et al. Submitted to *Transfusion*.