

Recent survey of Basque blood groups*

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J.N. Marshall Chalmers doktoreak 167 euskaldunen odola analizaturik, Rh negatiboak euskaldunen artean duen hedadura handiari buruz aurretik eginiko lanen ondorioak egiaztatu ditu.

Résultats des recherches du docteur J.N. Marshall Chalmers effectuées en Pays Basque sur un groupe de 167 Basques. Confirmant d'autres travaux antérieurs, le docteur Chalmers met en évidence la grande fréquence du Rh négatif chez les Basques.

El doctor J.N. Marshall Chalmers analizó a 167 vascos, confirmando las conclusiones de trabajos precedentes respecto a la gran extensión del Rh negativo entre los vascos.

* *Nature*, 1948, CLXX, p. 27.

We have determined the full *ABO*, *Rh* and *MN* blood groups of 167 Basques. The persons examined were selected as being of unmixed Basque descent, mostly by the Rev. Prof.

J. M. de Barandiaran. None of the persons examined was known to be a blood relation of any of the others. The accompanying tables give a summary of our results.

ABO Groups

| Group | Number | Frequency observed | Frequency expected | Gene frequencies |
|------------------|--------|--------------------|--------------------|-----------------------|
| O | 85 | 0.5090 | 0.5071 | A ₁ 0.2343 |
| A ₁ | 67 | 0.4012 | 0.4002 | A ₂ 0.0248 |
| A ₂ | 6 | 0.0359 | 0.0359 | B 0.0288 |
| B | 7 | 0.0419 | 0.0418 | 0 0.7121 |
| A ₁ B | 2 | 0.0120 | 0.0135 | |
| A ₂ B | 0 | 0.0000 | 0.0014 | |
| | 167 | 1.000 | 0.9999 | 1.0000 |

MN Groups

| Group | Number | Frequency observed | Frequency expected | Gene frequencies |
|-------|--------|--------------------|--------------------|------------------|
| M | 55 | 0.3293 | 0.3068 | M 0.5539 |
| MN | 75 | 0.4491 | 0.4942 | |
| N | 37 | 0.2216 | 0.1990 | N 0.4461 |
| | 167 | 1.0000 | 1.0000 | 1.0000 |

Rh Groups

| Reactions with antisera Anti-C D E c e | Rh phenotype frequencies Commonest genotype in each phenotype | Number | Frequency observed | Frequency expected |
|--|---|--------|--------------------|--------------------|
| - - - + + | cde/cde (rr) | 48 | 0.2874 | 0.3275 |
| - - - + + | CDe/cde(R,r) | 79 | 0.4731 | 0.3904 |
| + + + - - | CDe/CDe(R ¹ R ¹) | 13 | 0.0778 | 0.1246 |
| - + + + + | cDE/cde (R ² r) | 13 | 0.0778 | 0.0795 |
| + + + + + | CDe/eDE(R ¹ R ²) | 10 | 0.0599 | 0.0487 |
| + - - + + | Cde/ede (R ¹ r) | 3 | 0.0180 | 0.0180 |
| - + - + + | cDe/cde (R ² r) | 1 | 0.0060 | 0.0064 |
| - + + + - | cDE/cDE (R ² R ²) | 0 | 0.0000 | 0.0047 |
| + - - - + | Cde/Cde (R ¹ R ¹) | 0 | 0.0000 | 0.0002 |
| | | 167 | 1.0000 | 1.0000 |

Frequencies of gene combinations

| | |
|-----------------------|--------|
| cde (r) | 0.5723 |
| CDe (R ₁) | 0.3376 |
| cDE (R ₂) | 0.0688 |
| Cde (R ¹) | 0.0157 |
| cDe (R ₂) | 0.0056 |
| edE (R ¹) | 0.0000 |
| | 1.0000 |

The high frequency of *Rh*-negative (*D*-negative) persons (30.5 per cent) and of the *d* gene (58.8 per cent) are in agreement with the findings of Etcheverry¹ and support the views

recently put forward by one of us² regarding the history of the *d* gene. The low frequency of *cDE* is another notable feature. The excess of *Dd* heterozygotes and the deficiency of *MN* heterozygotes are clearly due to unavoidable errors of sampling. For this and other reasons steps are being taken to examine more persons. The combined results and a full discussion of them will be published elsewhere.

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1. Etcheverry, M.A., *Diá-méd.*, 17, 1237 (1945)

2. Mourant, A.E., *Nature*, 160, 505 (1947)