
From: Morgan, Timothy R (Long Beach) [mailto:timothy.morgan@va.gov]

Sent: Tuesday, October 09, 2007 8:05 PM

To: carringt@ncifcrf.gov

Cc: Bonkovsky, Herbert; Kristin Snow; Teresa Curto; dibiscam@slu.edu; everhartj@extra.niddk.nih.gov

Subject:

Mary

The HALT-C study group voted unanimously to permit you to send the HLA genotyping on HALT-C patients to your collaborators at Harvard and Microsoft, with the following understanding:

1. the results of their studies will be made freely available to the research community (i.e., the HLA data from HALT-C patients will not result in the private gain for a person, university, company or other entity),
2. racial categorization will be based on the racial categories used in the HALT-C study, and
3. your HLA genotyping dataset will be sent to NERI to be included in the HALT-C database.

Thanks again for your assistance with HLA testing of HALT-C specimens. We look forward to continued collaborations with you.

Tim Morgan and Herb Bonkovsky
For the HALT-C Steering Committee

Dear Colleagues:

As you know, we on the Genetics, Genomics, proteomics Committee have been working with Mary Carrington of NCI to obtain HLA typing on halt-c subjects. Mary has just written an email as follows:

“Zabrina Brumme in Bruce Walker's lab at Harvard and David Heckerman at Microsoft are trying to build a program to estimate 4 digit HLA typing from 2 digit typing, which will be really useful to the scientific community. They have asked me for 4 digit class I typing on as many samples as possible. They do not need any identifier whatsoever, no clinical data, or anything else except for the HLA type and race. I am busy asking all cohorts, all of whom have said fine so far. Would it be okay to send them typings from HALT-C? They could list HALT-C in the acknowledgements if you like. We would need race on the people we have typed to make the data useful to them. We could send a list of the typings to whoever could help us with this. Many thanks for letting me know.
Best,
Mary.”