## CCSS Genetics Working

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#### Current proposals from baseline

- Single gene diseases describing phenotype in the cohort
- LFS/LFL family history
  - 44 families
- Colon cancer history in first degree relatives
  - 139 families
  - 13 families with colon cancer and sibling with cancer
  - 6 families with sibling with colon cancer
  - All other families have colon cancer in parents and no cancers in any full sibs.
  - Proband cancers include leukemia, HD, NHL, CNS, Bone and STS, Wilms and NBL.
  - Some suggest Turcot or Lynch.
  - 25 have occurred in a parent under the age of 45 years

#### Current proposals using biospecimens

- MDM2 (SNP309) analysis in Hodgkin lymphoma survivors treated with radiotherapy +/- breast cancer
- Recommendation: Replenish depleted biospecimens and consider increasing repository

### Future proposals

- GWA combined with FH for NMSC
- GWA for Hodgkin lymphoma
  - Two stage design: Discovery; Replication
  - Primary aim etiology
  - Secondary aim SMN (or other adverse outcome with sufficient prevalence)
  - Potentially include incident cases from COG HD
    - Compare genotypes that might predict outcome or late effects in the COG group based on the CCSS group data

#### Priorities for 2007 survey

- Develop web-based family history survey
  - Present baseline history
  - Probing questions for ambiguous answers
  - Ask for corrections and updates: FH of cancer, new relatives (including offspring)
- Target high yield populations first
  - Hodgkin lymphoma SMNs
  - Sarcoma known genetic predisposition; SMNs
  - All SMNs (w/controls matched by latency and primary diagnosis)

**Questions: IT platform?** 

Can we update FH on deceased probands?

#### Expanded Cohort

- Web-based family history
- Hypothesis that siblings, offspring with cancer and probands with SMNs enrich FH data and may form sub-population with genetic cancer predisposition
- Inform methodology from 2007 survey
- Biosample collection

# Additional recommendations: Biospecimens

- Biospecimen collection
  - Existing cohort: Re-fill depleted samples
  - Expanded cohort: Early specimen collection
- Advisory committee to be established to evaluate proposals using biospecimens followed by recommendations of biospecimens to genetics working group and other CCSS working groups
- Material Transfer Agreement for biospecimens

#### Public Use Database

#### Concerns

- Inaccuracy of data not validated or clarified
- Potential for identification of probands based on family history
- Could family history be provided by request?