WashU Epigenome Browser - Feature # 161

| Status: | New | Priority: | Normal |
|--|--|---|--|
| Author: | Rebecca Lowdon | Category: | |
| Created: | 02/13/2014 | Assignee: | Xin Zhou |
| Updated: | 02/13/2014 | Due date: | |
| Subject: | Include scaffolds on genome | snapshot | |
| Description | | | |
| | - | | referred to as "NA###") in the "genome snapshot like you do now, but have a button to |
| snapshot" app. M | aybe make it an optional feature ge | enerate the "standard" genome | |
| snapshot" app. M generate addition You can get to the | aybe make it an optional feature ge al snapshot over the scaffolds and un | enerate the "standard" genome assigned chromosome fragme nes it's nice to relocate from the | snapshot like you do now, but have a button to |
| snapshot" app. M generate addition You can get to the | aybe make it an optional feature ge al snapshot over the scaffolds and un ese fragments by search, but sometin | enerate the "standard" genome assigned chromosome fragme nes it's nice to relocate from the | snapshot like you do now, but have a button to onts (most interested in scaffolds at this point). |
| snapshot" app. M generate addition You can get to the | aybe make it an optional feature ge al snapshot over the scaffolds and un ese fragments by search, but sometin | enerate the "standard" genome assigned chromosome fragme nes it's nice to relocate from the | snapshot like you do now, but have a button to onts (most interested in scaffolds at this point). |
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