Today's Meeting

Hoch,Jeffrey <hoch@uchc.edu>   Tue, Mar 7, 2017 at 3:42 PM
To: Joel Markus <joel@jmarkusdesign.com>, "Schuyler,Adam" <schuyler@uchc.edu>
Cc: Visalli Media <gina@visallimedia.com>

Thanks Joel!

Below is a first stab at categories and exemplars.

- J

1. Spectral

    a. Processing

       NMRPipe

       RNMRTK

       NMRfx

    b. Visualization

       NMRDraw

       Sparky

       NMRViewJ

2. Assignment

   PINE

   Abacus

3. Relaxation

   ModelFree

   DASHA

4. Chemical shift

    a. Prediction

        SHIFTS

        SHIFTX

        SHIFTX2

    b. Analysis

        TALOS+

        CS23D

        CS-ROSETTA

5. Molecular modeling

    a. Dynamics simulation

        AMBER

        GROMACS

        MOSART

    b. Docking

        HADDOCK

    c. Strucure determination

        AMBER

        XPLOR-NIH

        CS-ROSETTA

6. Structure

    a. Visualization

        MOLMOL

        VMD

        PyMOL

    b. Validation

ASDP

ProCheckNMR

7. Redidual dipole couplings

REDCAT

REDCRAFT

PALES

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From: Joel Markus <joel@jmarkusdesign.com>
Sent: Tuesday, March 7, 2017 3:12 PM
To: Schuyler,Adam; Hoch,Jeffrey
Cc: Visalli Media
Subject: Today's Meeting

Adam and Jeff,

Thanks for taking the time to meet this morning to go over a few fundamental flow diagrams for content flow and navigation.

Attached are the two documents that were presented this morning.