| Week | Lecture | Date | Content | HW Out | t HW In |
|------|---------|------|-------------------------------------|---------|---------|
| 1 | 1 | 1/17 | introduction to protein engineering | | |
| | 2 | 1/19 | primary structure | | |
| 2 | 3 | 1/22 | primary structure | | |
| | 4 | 1/24 | secondary structure | | |
| | 5 | 1/26 | secondary structure | | |
| 3 | 6 | 1/29 | secondary structure | HW 1 | |
| | 7 | 1/31 | motif | | |
| | 8 | 2/2 | tertiary structure | | |
| 4 | 9 | 2/5 | tertiary structure | HW 2 | HW 1 |
| | 10 | 2/7 | tertiary structure | | |
| | 11 | 2/9 | quaternary structure | | |
| 5 | 12 | 2/12 | protein core | | HW 2 |
| | 13 | 2/14 | protein core | | |
| | 14 | 2/16 | | Mid Ter | m 1 |
| 6 | 15 | 2/19 | amino acid alphabet | | |
| | 16 | 2/21 | thermophilic proteins | | |
| | 17 | 2/23 | thermophilic proteins | | |
| 7 | 18 | 2/26 | interface | | |
| | 19 | 2/28 | interface | | |
| | 20 | 3/2 | 2nd, 3rd structure prediction | HW 3 | |
| 8 | 21 | 3/5 | knowledge-based design | | |
| | 22 | 3/7 | knowledge-based design | | |
| | 23 | 3/9 | transmembrane protein | | HW 3 |
| 9 | 24 | 3/19 | forcefield | | |
| | 25 | 3/21 | rotamer | HW 4 | |
| | 26 | 3/23 | search algorithm | | |
| 10 | 27 | 3/26 | computational protein design | | |
| | 28 | 3/28 | computational protein design | | |
| | 29 | 3/30 | ТВА | | HW 4 |
| 11 | 30 | 4/2 | ТВА | | |
| | 31 | 4/4 | bioinformatics | | |
| | 32 | 4/6 | | Mid Ter | m 2 |
| 12 | 33 | 4/9 | directed evolution | | |
| | 34 | 4/11 | directed evolution | | |
| | 35 | 4/13 | directed evolution | | |
| 13 | 36 | 4/16 | protease | | |
| | 37 | 4/18 | enzyme | HW #5 | |
| | 38 | 4/20 | carbohydrate binding protein | | |
| 14 | 39 | 4/23 | DNA binding protein | | |
| | 40 | 4/25 | ТВА | | |
| | 41 | 4/27 | biomaterial | | HW #5 |
| 15 | 42 | 4/30 | modeling | | |