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r"""
Updates to the thin SVD using NumPy.
This function is a SAGE replication of Matthew Brand's article on "Fast low-rank modifications of the thin singular value decomposition." <a href="http://www.stat.osu.edu/-dmsl/thinSVOtracking.pdf">http://www.stat.osu.edu/-dmsl/thinSVOtracking.pdf</a>. This function is an approximation to the true thin SVO, therefore, no tests are provided.
AUTHORS:
- Taylor Steiger, James Pak (2013-06-10): initial version
EXAMPLES::
        \begin{aligned} &\text{sage: } X = \text{np.array}([[1.0,2.0,3.0,4.0],[3.0,2.0,5.0,5.0],[5.0,3.0,1.0,1.0],[7.0,7.0,7.0,7.0]])} \\ &\text{sage: } U, \text{s, } V = \text{np.linolg.svd}(X, \text{full_matrices} = \text{Folse}) \\ &\text{sage: } a = \text{np.reshpae(pn.array([4.0,5.0,1.0,7.0]), (-1, 1))} \\ &\text{sage: } U, \text{s, } V = \text{svd\_update(U, np.diag(S), V, X, a, update} = \text{True}) \end{aligned} 
       Downdate
        soge: X = np.arroy([[1.0,2.0,3.0,4.0],[3.0,2.0,5.0,5.0],[5.0,3.0,1.0],[7.0,7.0,7.0,7.0]])
soge: U, s, V = np.lindg.svd(X, full_matrices = False)
soge: U, S, V = svd.update(U, np.diag(S), V, X, domadate = True)
        \label{eq:sog: X = np.array([[1.0,2.0,3.0,4.0],[3.0,2.0,5.0,5.0],[5.0,3.0,1.0,1.0],[7.0,7.0,7.0,7.0]])} \\ soge: U, s, V = np.lindg.svd(X, full_matrices = False) \\ soge: 0 = np.respace(np.array([4.0,5.0,1.0,7.0]), (-1, 1)) \\ soge: U, S, V = svd.update(U, np.diag(s), V, X, o) \\ \\
       Recenter
       \begin{split} & \text{sage: } X = \text{np.array}([[1.0, 2.0, 3.0, 4.0], [3.0, 2.0, 5.0, 5.0], [5.0, 3.0, 1.0, 1.0], [7.0, 7.0, 7.0]]) \\ & \text{sage: } U, \text{ s, } V = \text{np.linalg.svd}(X, \text{ full_matrices} = \text{False}) \\ & \text{sage: } U, \text{ S, } V = \text{svd\_update}(U, \text{ np.diag}(\text{s}), \text{ V, } X) \end{split}
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import numpy as np
def svd_update(U, S, V, X, c = None, update = False, downdate = False):
      INPUT:
       - U -- a (nxn) matrix containing singular vectors of \mathbf{X}.
      - S -- a (nxn) diagonal matrix containing singular values. the ith diagonal entry is the singular value corresponding to the ith column of U
       - V -- a (nxn) matrix containing singular vectors of X.
       - X -- a (mxn or nxm) matrix such that U^T*X*V=S.
       - c -- (default: None) a column vector for revision or update of decomposition.
       - update -- (default: False) boolean whether to add c to the decomposition. If true, c must also be provided.
       - downdate -- (default: False) boolean whether to downdate the decomposition.
       OUTPUT:
       A 3-tuple consisting of matrices in this order:

    Transformed U.
    Transformed S.
    Transformed V.

        The SVD rank-1 modification algorithm is described by Matthew Brand in the paper at, http://www.stat.osu.edu/~dmsl/thinSVDtracking.pdf>
The algorithm works as follows:
      During testing, this code showed strong deviations, since it is an approximation, from the true thin SVD of the matrix X. However, testing was conducted with a small matrix X, and may be working perfectly fine. ^{\rm max}
        V = np.vstack([V. np.zeros(V.shape[1])])
        v = np.vstack([v, np.zeros(v.snape[1])])
if down or type(c) == type(np.array([])):
    b = np.zeros(V.shape[0])
    b[-1] = 1
    b = np.reshape(b, (b.shape[0], 1))
             \begin{split} & \text{m} = \text{np.reshape(np.dot(np.transpose(U), a), (-1, 1))} \\ & p = \text{np.reshape(a - np.dot(U, m), (-1, 1))} \\ & \text{Re = np.linal, norm(p)} \\ & P = \text{np. reshape(np.multiply((1 / Ra), p), (-1, 1))} \\ & P = \text{np. reshape(np.multiply((1 / Ra), p), (-1, 1))} \\ & \text{q} = \text{b} - \text{np.dot(V, n)} \\ & \text{q} = \text{b} - \text{np.dot(V, n)} \\ & \text{B} = \text{np.linal, norm(q)} \\ & \text{Q} = \text{np.reshape(np.multiply((1 / Rb), q), (-1, 1))} \end{split}
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\label{eq:continuous_series} \begin{split} k &= S \\ K &= np.zeros((k.shape[\emptyset] + 1, k.shape[\emptyset] + 1)) \\ K[:-1,:-1] &= k \\ stock &= np. vstock(np.append(m, R0)) \\ t &= np.reshape(np.append(n, R0), (1, -1)) \\ dot &= np.dot(stock, t) \\ K &= np.add(K, dot) \end{split}
D, P = np.linalg.eig(K)
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return (np.transpose(np.linalg.inv(P)), np.diag(D), P)