

Calcul Scientifique

Projet de Calcul Scientifique

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1.

1.1.

Matrix dimension	Matrix type	Exec. time for eig (s)	Exec. time for power_v11, (s)
200×200	Type 1	9.000e-02	1.510e+00
400×400	Type 1	4.000e-02	1.831e + 01
600×600	Type 1	6.000e-02	6.021e+01
200×200	Type 2	3.000e-02	3.000e-02
400×400	Type 2	4.000e-02	4.000e-02
600×600	Type 2	7.000e-02	1.700e-01
200×200	Type 3	1.000e-02	5.000e-02
400×400	Type 3	3.000e-02	5.200e-01
600×600	Type 3	7.000e-02	1.270e+00
200×200	Type 4	2.000e-02	1.670e + 00
400×400	Type 4	3.000e-02	2.094e+01
600×600	Type 4	6.000e-02	5.456e + 01

Table 1: Execution time for different sizes and types of matrices

We can see that the power_v11 algorithm is generally slower than the eigen function especially for the type 2 and 4 matrices.

1.2.

```
nb_it = 1;
norme = norm(beta*v - z, 2)/norm(beta,2);

while(norme > eps && nb_it < maxit)
    beta_old = beta;
    v = z/norm(z, 2);
    z = A*v;
    beta = (v'*z)/(v'*v);
    norme = abs(beta-beta_old)/abs(beta_old);
    nb_it = nb_it + 1;
end</pre>
```

Listing 1: Inner loop of the new algorithm

Matrix dimension	Matrix type	Exec. time for $power_v11$, (s)	Exec. time for $power_v12$, (s)	
200×200	Type 1	1.960e+00	3.200e-01	
400×400	Type 1	1.888e+01	2.660e+00	
600×600	Type 1	5.031e+01	7.070e+00	
200×200	Type 2	1.000e-02	1.000e-02	
400×400	Type 2	7.000e-02	1.000e-02	
600×600	Type 2	1.800e-01	4.000e-02	
200×200	Type 3	3.000e-02	1.000e-02	
400×400	Type 3	6.100e-01	1.100e-01	
600×600	Type 3	1.270e+00	2.600e-01	

200×200	Type 4	1.530e+00	2.900e-01
400×400	Type 4	2.113e+01	3.060e+00
600×600	Type 4	5.914 + e01	6.480e+00

We can see that the power_v12 algorithm is globally faster than the power_v11.

1.3.

The main drawback of the deflated power method is the numerous matrix-vector products required to compute the eigenvectors as well as the fact that each iteration compute only one eigenvalue which can be slow if a lot of eigenvalues are desired.

1.4.

If we apply Algorithm 1 to m vectors, there is no reason for the columns of V to converge to a base. Each vector will converge toward a different projection of the dominant eigenvalue.

1.5.

In Algorithm 2, the matrix H is a smaller matrix, with dimension $n \times m$, therefore, even for larger matrices A, computing the spectral decomposition of H will not be computionally expensive.

1.6.

1.7.

- 1: function Subspace iter v1 (Raleigh-Ritz Projection)
- 2: Input : $A \in \mathbb{R}^{n \times n}$, ε , MaxIter, PercentTrace
- 3: Output : $n_{\rm ev}$ dominant eigenvectors $V_{\rm out}$ and the corresponding eigenvalues $\Lambda_{\rm out}$
- 4: Generate an initial set of m orthonormal vectors $V \in \mathbb{R}^{n \times m}$; k = 0; PercentReached = 0
- 5: repeat until Percent Reached > Percent Trace $\lor n_{\mathrm{ev}} = m \lor k > \text{MaxIter}$
- 6: $k \leftarrow k+1$
- 7: Compute Y such that $Y = A \cdot V$
- 8: $V \leftarrow$ orthonormalisation of the columns of Y
- 9: Rayleigh-Ritz projection applied on matrix A and orthonormal vectors V
- 10: Convergence analysis step: save eigenpairs that converged and update PercentReached

1.8.

1.9.

1.10.

Matrix dimen-	Matrix type	Flops for	Flops for	Flops for	$p(A^p)$
sion		subspace_iter0	subspace_iter1	subspace_iter2	
200×200	Type 1	2309	263	132	2
200×200	Type 1	2309	263	88	3
200×200	Type 1	2309	263	53	5
200×200	Type 1	2309	263	27	10

When increasing the valu of p to compute A^p in subspace_iter2, the number of flops to compute the results is: Flops(iter2) $\simeq \frac{\text{Flops}(\text{iter2})}{n}$.

1.11.

1.12.

By freezing the converged columns, the algorithm will not have to recalculate them everytime. Which means that the accuracy for the eigenpairs will be more equal. The first and last will have the same approximate size.

2.

2.1.

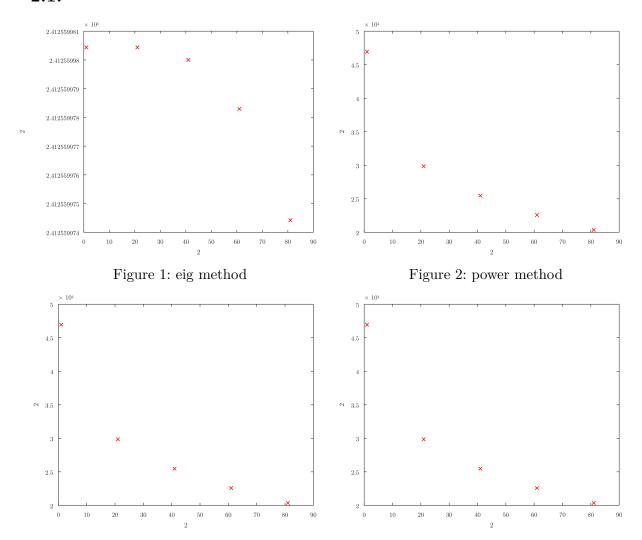


Figure 3: subspace_iter0 method

Figure 4: subspace_iter1 method

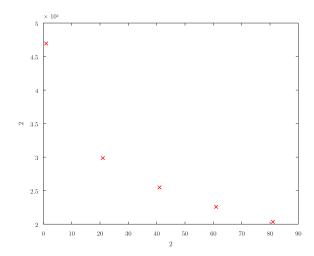


Figure 5: subspace_iter2 method